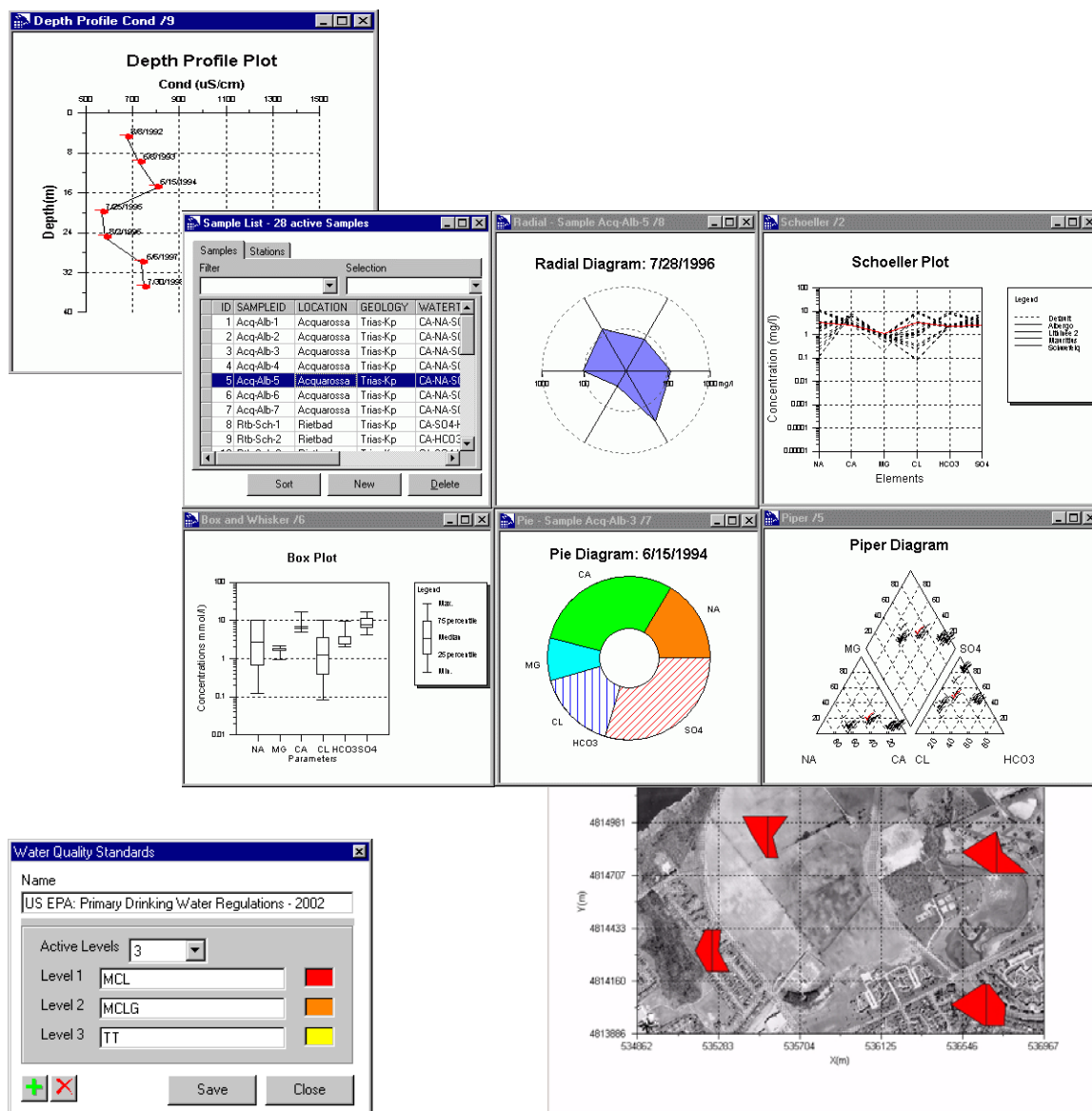


AquaChem 2011.1

User's Manual

Water Quality Data Analysis, Plotting, and Modeling



Preface

Schlumberger Water Services (SWS) is a recognized leader in the development and application of innovative groundwater technologies in addition to offering expert services and professional training to meet the advancing technological requirements of today's groundwater and environmental professionals.

Schlumberger Water Services software consists of a complete suite of environmental software applications engineered for data management and analysis, modeling and simulation, visualization, and reporting. Schlumberger Water Services software is currently developed by SWS and sold globally as a suite of desktop solutions.

For over 18 years, our products and services have been used by firms, regulatory agencies, and educational institutions around the world. We develop each product to maximize productivity and minimize the complexities associated with groundwater and environmental projects. To date, we have over 14,000 registered software installations in more than 85 countries!

Need more information?

If you would like to contact us with comments or suggestions, you can reach us at:

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Obtaining Technical Support

When contacting the Technical Support department, please have the following information available:

- your Name, Company Name, and Address
- your Product Serial Number, Version, and Build number
- full pathname to your software installation, and to your program files

You may send us your questions via e-mail, fax, or call one of our technical support specialists. Please allow up to two business days for a response. Technical support is available 6:00 am to 6:00 pm EST Monday to Friday (excluding Canadian holidays).

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Schlumberger Water Services also offers expert consulting and peer reviewing services for data management, groundwater modeling, aqueous geochemical analysis, and pumping test analysis. For further information, please contact sws-services@slb.com.

Schlumberger Water Services Software

We also develop and distribute a number of other useful software products for the groundwater professionals, all designed to increase your efficiency and enhance your technical capability, including:

- Visual MODFLOW Premium*
- Hydro GeoBuilder*
- Hydro GeoAnalyst*
- Aquifer Test Pro*
- AquaChem*
- GW Contour*
- UnSat Suite Plus*
- Visual HELP*
- Visual PEST-ASP

Visual MODFLOW Premium

Visual MODFLOW Premium is a three-dimensional groundwater flow and contaminant transport modeling application that integrates MODFLOW-2000, SEAWAT, MODPATH, MT3DMS, MT3D99, RT3D, VMOD 3D-Explorer, WinPEST, Stream Routing Package, Zone Budget, MGO, SAMG, and PHT3D. Applications include well head capture zone delineation, pumping well optimization, aquifer storage and recovery, groundwater remediation design, simulating natural attenuation, and saltwater intrusion.

Hydro GeoBuilder

Hydro GeoBuilder provides a flexible and streamlined approach to developing conceptual models for Visual MODFLOW and FEFLOWTM. Featuring a powerful multi-format object/data import tool, Hydro GeoBuilder offers modeling professionals an expanded workbench of two- and three- dimensional tools

for conceptualizing the hydrogeologic environment, in addition to increased flexibility for assigning model properties independent of the finite difference grid or finite element mesh. This means you save hours when building your numeric model.

FEFLOW is a registered trademark of DHI-WASY

Hydro GeoAnalyst

Hydro GeoAnalyst is an information management system for managing groundwater and environmental data. Hydro GeoAnalyst combines numerous pre and post processing components into a single program. Components include, Project Wizard, Universal Data Transfer System, Template Manager, Materials Specification Editor, Query Builder, QA/QC Reporter, Map Manager, Cross-Section Editor, HGA 3D-Explorer, Borehole Log Plotter, and Report Editor. The seamless integration of these tools provide the means for compiling and normalizing field data, analyzing and reporting subsurface data, mapping and assessing spatial information, and reporting site data.

AquiferTest Pro

AquiferTest Pro, designed for graphical analysis and reporting of pumping test and slug test data, offers the tools necessary to calculate an aquifer's hydraulic properties such as hydraulic conductivity, transmissivity, and storativity. AquiferTest Pro is versatile enough to consider confined aquifers, unconfined aquifers, leaky aquifers, and fractured rock aquifers conditions. Analysis results are displayed in report format, or may be exported into graphical formats for use in presentations. AquiferTest Pro also provides the tools for trends corrections, and graphical contouring water table drawdown around the pumping well.

AquaChem

AquaChem is designed for the management, analysis, and reporting of water quality data. AquaChem's analysis capabilities cover a wide range of functions and calculations frequently used for analyzing, interpreting and comparing water quality data. AquaChem includes a comprehensive selection of commonly used plotting techniques to represent the chemical characteristics of aqueous geochemical and water quality data, as well includes PHREEQC - a powerful geochemical reaction model.

GW Contour

The GW Contour data interpolation and contouring program incorporates techniques for mapping velocity vectors and particle tracks. GW Contour incorporates the most commonly used 2D data interpolation techniques for the groundwater and environmental industry including Natural Neighbor, Inverse Distance, Kriging, and Bilinear. GW Contour is designed for contouring surface or water levels, contaminant concentrations, or other spatial data.

UnSat Suite Plus

UnSat Suite Plus seamlessly integrates multiple one-dimensional unsaturated zone flow and solute transport models into a single, intuitive working environment. Models include SESOIL, VS2DT, VLEACH, PESTAN, Visual HELP and the International Weather Generator. The combination of models offers users the ability for simulating the downward vertical flow of water and the migration of dissolved contaminants through the vadose zone. UnSat Suite Plus includes tools for project management, generating synthetic weather data, modeling flow and contaminants through the unsaturated zone, estimating groundwater recharge and contaminant loading rates, and preparing compliance reports.

Visual HELP

Visual HELP is a one-dimensional, unsaturated zone flow modeling application built for optimizing the hydrologic design of municipal landfills. Visual HELP is based on the US E.P.A . HELP model (Hydrologic Evaluation of Landfill Performance) and has been integrated into a 32-Bit Windows application. It combines the International Weather Generator, Landfill Profile Designer, and Report Editor. Applications include designing landfill profiles, predicting leachate mounding, and evaluating potential leachate seepage to the groundwater.

Visual PEST-ASP

Visual PEST-ASP combines the powerful parameter estimation capabilities of PEST-ASP, with the graphical processing and display features of WinPEST. Visual PEST-ASP can be used to assist in data interpretation, model calibration and predictive analysis by optimizing model parameters to fit a set of observations. This popular estimation package achieves model independence through its capacity to communicate with a model through its input and output files.

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1

Introduction to AquaChem

AquaChem is a software package developed specifically for graphical and numerical analysis and modeling of water quality data. It features a fully customizable database of physical and chemical parameters and provides a comprehensive selection of analysis tools, calculations, and graphs for interpreting water quality data.

AquaChem's data analysis capabilities cover a wide range of functionalities and calculations including unit conversions, charge balances, sample comparison and mixing, statistical summaries, trend analyses, and much more. AquaChem also comprises a customizable database of water quality standards with up to three different action levels for each parameter. Any samples exceeding the selected standard are automatically highlighted with the appropriate action level color for easily identifying and qualifying potential problems.

These powerful analytical capabilities are complemented by a comprehensive selection of commonly used plotting techniques to represent the chemical characteristics of water quality data. The plot types available in AquaChem include:

- Correlation plots: X-Y Scatter, Ludwig-Langelier, Wilcox, Depth Profile
- Summary plots: Box and Whisker (Grouped, Multiple Parameters, Multiple Stations, Time), Frequency Histogram, Quantile, Detection summary, Meteoric Water Line, Stacked Bars
- Multiple parameter plots: Piper, Durov, Ternary, Schoeller
- Time-Series plots (multiple parameters, multiple stations, statistical)
- Geothermometer and Giggenbach plot
- Detection summary plot
- Probability plots
- Single sample plots: Radial, Stiff, and Pie
- Thematic Map plots: Bubble, Pie, Radial and Stiff plots at sample locations

Each of these plots provides a specific interpretation of the many complex interactions between the groundwater and aquifer materials, and identifies important data trends and groupings.

In addition, AquaChem features a built-in link to the popular geochemical modeling program PHREEQC for calculating equilibrium concentrations (or activities) of

chemical species in solution and saturation indices of solid phases in equilibrium with a solution. For more advanced simulations, you may link to the USGS programs PHREEQCI or PHREEQC for Windows, and use your AquaChem samples as input solutions for these modeling utilities.

Once you start using AquaChem, you will see that it is truly one of the most powerful tools available for interpretation, analysis and modeling of any water quality data set.

1.1 Installing AquaChem

1.1.1 System Requirements

AquaChem requires the following **minimum** hardware configuration:

Supported Operating Systems:

Windows 7 Professional, Enterprise or Ultimate

Windows Vista Business, Ultimate or Enterprise

Windows XP Pro (SP3)

Note: Home and Starter Versions are not supported.

Processor: Pentium 4, 32-bit or 64-bit

CD or DVD Drive: CD-ROM Drive

Hard Disk: 100MB

RAM: 512 MB (1GB or more Recommended)

Networking Hardware: Network Card (required for licensing)

The AquaChem installation package requires the following software configuration:

Microsoft .NET Framework 2.0 or higher

1.1.2 Installation

Stand Alone Installation

AquaChem is distributed on one CD-ROM. To install, please follow these directions:

- Place the CD into your CD-ROM drive and the initial installation screen should load automatically. Once loaded, an installation interface with several different tabs will be presented.
- Please take the time to explore the installation interface, as there is information concerning other Schlumberger Water Services products, our worldwide distributors, technical support, consulting, training, and how to contact us.
- On the initial Installation tab, you may choose from the following two buttons: **AquaChem Installation** and **AquaChem User's Manual**
- The User's Manual button will display a PDF document of the manual, which requires the Adobe Reader to view. If you do not have the Adobe Reader, a link has been created in the interface to download the appropriate software.

- The Installation button will initiate the installation of AquaChem on your computer. AquaChem must be installed on your local hard disk in order to run. Follow the installation instructions, and read the on-screen directions carefully. You will be prompted to enter your name, company name and serial number. Please ensure that you enter your serial number exactly as it appears on your CD case or invoice. Be sure to use capital letters and hyphens in the correct locations.
- Once the installation is completed, you must re-boot your computer for the system changes to take effect. After the installation is complete and your system has re-booted, you should see the blue AquaChem icon on your Desktop screen labeled **AquaChem 2011.1**. To start working with AquaChem, double-click on this icon.

To install the software from the CD-ROM without the aid of the installation interface, you can:

- Open Windows Explorer, and navigate to the CD-ROM drive
- Open the Installation folder
- Double-click the **AquaChem_Setup.MSI** to initiate the installation

Follow the on-screen installation instructions, which will lead you through the install and subsequently produce a desktop icon for you.

PHREEQC-I Installation

The USGS's PHREEQC-Interactive program is a graphical interface for preparing and running complex geochemical modeling scenarios. AquaChem has a built-in link to the PHREEQC-Interactive program that is capable of launching this program with all selected samples already formatted as modelling input.

The PHREEQC-I must be installed separately; the installation file is available on your CD-ROM under the **PHREEQC** directory.

Once installed, the PHREEQCI executable must be registered in the Aquachem preferences. It can then be launched from AquaChem (Tools / Modeling / PHREEQC (Advanced)) and the input file will automatically be initialized with the chemical composition of the samples that are highlighted in the AquaChem sample list.

PHREEQC for Windows Installation

AquaChem also supports a link to the PHREEQC for Windows program. This program is an alternative graphical interface that also allows for preparing and running advanced geochemical simulations.

If you wish to install and use the PHREEQC for Windows program, the installation is available in the **PHREEQC** folder on the installation CD-ROM.

These files are also available for download from the Schlumberger Water Services FTP site:

<ftp.flowpath.com/software/aquachem>

and from the USGS - PHREEQC home page:

http://wwwbrr.cr.usgs.gov/projects/GWC_coupled/phreeqc/

1.2 Uninstalling AquaChem

There may be instances where you will need to uninstall (remove) AquaChem from your system (i.e. if the software is to be transferred to another computer, or you need to reinstall on the current computer). To uninstall AquaChem:

- Locate the **Add/Remove Programs** option in your **Windows' Control Panel**.
- Select **AquaChem 2011.1** as the program to be removed
- Follow the on-screen instructions.
- Once you are finished, re-boot your system to ensure all system files are updated.

1.3 On-Line Help

This manual is available in two forms: as a printed book, and as an online help file. To view the online help version of this manual, select **Help > Contents**.

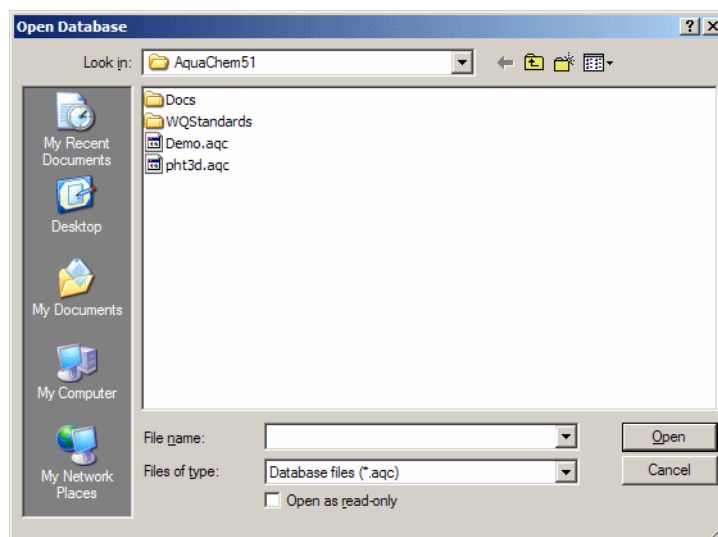
1.4 Starting AquaChem



To start AquaChem, you must have the program installed on your hard disk. If you have not yet installed AquaChem, please refer to the section, Installing AquaChem, which is described above. Otherwise start AquaChem by double-clicking on the desktop icon (as shown on the left-hand side), or by accessing

SWS Software/Aquachem 2011.1 from your **Start > Programs** Windows menu.

Upon starting AquaChem, the following **Open Database** dialogue will be displayed prompting you to select a valid AquaChem database.



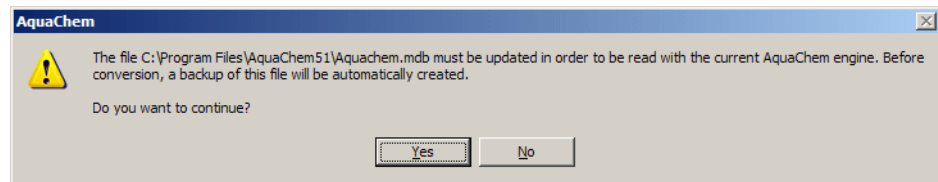
Select the **Demo.aqc** file to open the demonstration database; to open a different database, browse to the appropriate folder. Otherwise, to create a new database click **[Cancel]** in this dialogue and select **File > New** from the main menu.

1.4.1 Opening Old Project Sets

You may open an .AQC file from previous versions of AquaChem.

NOTE: AquaChem no longer supports direct opening of 3.x projects and the .HC3 extension has been removed from AquaChem Open File interface. To open a 3.x project in AquaChem you have to export the data into a text, MS Access, or MS Excel file and import it into a new database created in the current version of AquaChem.

To open a project from v.4.0 or higher, use the **File > Open** command. You will be prompted with an **Open Database** dialogue. Browse to the folder which contains your database and press **[Open]**. The following message will appear.



If you select **[Yes]**, the AquaChem database will be opened with a screen layout as shown on the next page, or if you select **[No]** then the option of opening the old project sets will be canceled.

1.5 AquaChem Interface Layout

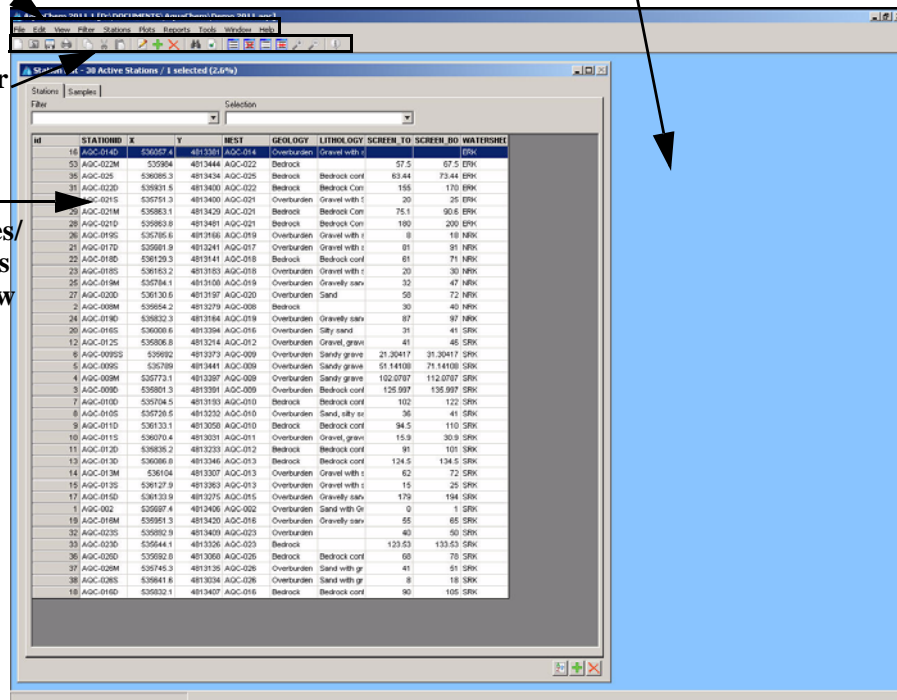
After opening an AquaChem database file, a screen layout similar to the following figure will appear.

Main Menu Bar

Parent Window

**Main
Toolbar**

**Active
Samples/
Stations
Window**



Parent Window is the main AquaChem window which houses all other windows.

Main Menu Bar contains specific menus for graphs and dataset. Depending upon the currently selected window, each window has a distinct set of menu options. A detailed description of each main menu options associated with various windows is provided in Chapter 3: AquaChem Menu Commands.

Main Toolbar contains specific tool buttons for different options. A detailed description of each main toolbar item is provided in section 1.6 of this chapter, AquaChem Toolbar.

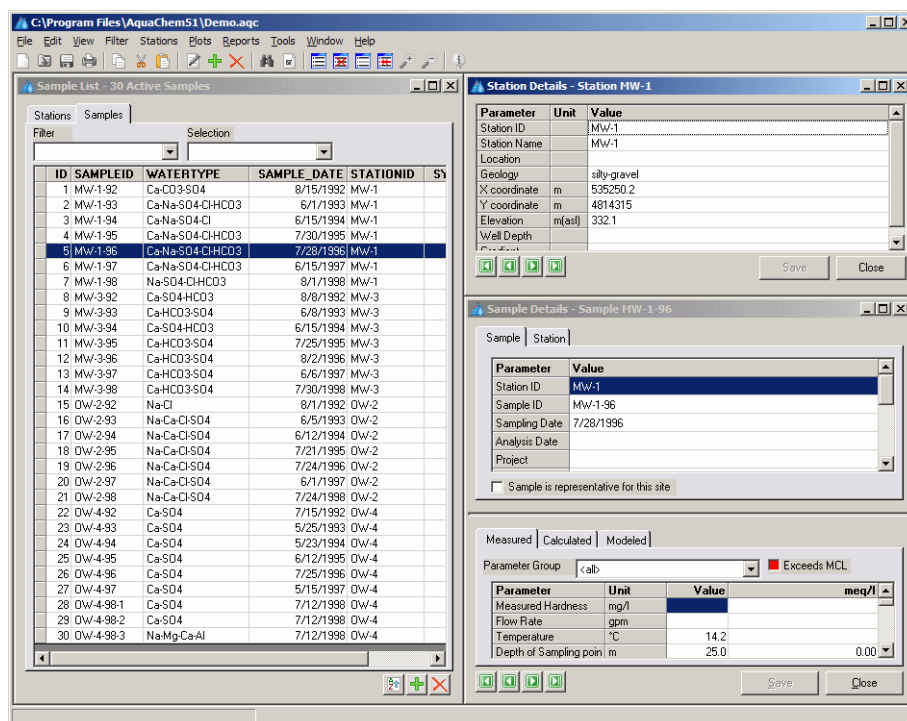
Active Samples/Stations Window will always appear when you open an AquaChem database and will remain on-screen as long as the project database is open (i.e. the Active Samples/Stations window cannot be closed unless the project database is closed). This window displays the list of samples and stations in the currently selected database. Two further windows can be accessed through the **Active Samples/Stations** tab to display and manipulate the dataset:

- **Sample Details Window** contains details for the selected sample.
- **Station Details Window** contains details for the selected station.

The following remaining 'Child' windows are used to display and manipulate the data which can be accessed through the main menu commands:

- **Table View** available under **View** menu allows you to view and edit the data in the database as a table.
- **Template Designer** available under the **File** menu contains options for designing print templates for plots and reports.
- **Reports** loads pre-defined data analysis reports, or user-designed reports. The **Report Designer** available under the **Reports** contains options for designing data reports.
- **Tools** loads several tools for data analysis and interpretation. **Modeling > PHREEQC** available under the **Tools** loads the interface for the PHREEQC modeling utility, and provides direct links to the graphical user interfaces for PHREEQC-I and PHREEQC for Windows.

AquaChem follows most standard Windows interface conventions. Each window can be minimized to the bottom of the Parent window and re-opened as needed. Likewise, window sizes can be adjusted by dragging and releasing the corners of the window frame. Windows can be arranged (as shown below for example) on the Parent window using the **Windows > Tile Horizontal** or **Tile Vertical** command which are available from the main menu.



The following section summarizes the features of each of the main AquaChem windows.

1.5.1 Active Samples/Stations Window

AquaChem follows a database hierarchy of stations followed by samples. This means that each sample must have a corresponding station. When you create a new sample, a corresponding station must be assigned to it.

The **Active Sample/Stations** window contains summarized information about every active sample and station in the database; the fields in this window are read-only which means that fields in this window cannot be edited. This window contains two tabs: **Stations** and **Samples**. Clicking on these tabs displays the following windows.

Station List - 4 Active Stations / 0 selected (0.0%)

Station Samples Selection

Filter

ID	STATION	X	Y	GEOLOGY	ELEVATION
1	MW-1	535250.2	4814315	silty-gravel	332.1
3	MW-2	535535.5	4814905	silty-sand	333.9
2	MW-3	536688.1	4814036	sandy-silt	334.8
4	MW-4	536720.7	4814826	sandy-grave	335.4

Sample List - 26 Active Samples / 0 selected (0.0%)

Stations Samples Selection

Filter

ID	Station	Date	Code	WATERTY	GEOLOGY	SYMBOL	REP
1	MW-1	8/15/1992	MW-1-92	Ca-Na-SO4	silty-gravel	18	<input type="checkbox"/>
2	MW-1	6/1/1993	MW-1-93	Ca-Na-SO4	silty-gravel	18	<input type="checkbox"/>
3	MW-1	6/15/1994	MW-1-94	Ca-Na-SO4	silty-gravel	18	<input checked="" type="checkbox"/>
4	MW-1	7/30/1995	MW-1-95	Ca-Na-SO4	silty-gravel	18	<input type="checkbox"/>
5	MW-1	7/28/1996	MW-1-96	Ca-Na-SO4	silty-gravel	18	<input type="checkbox"/>
6	MW-1	6/15/1997	MW-1-97	Ca-Na-SO4	silty-gravel	18	<input type="checkbox"/>
7	MW-1	8/1/1998	MW-1-98	Na-SO4-Cl-H	silty-gravel	18	<input type="checkbox"/>
15	MW-2	8/1/1992	DW-2-92	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
16	MW-2	6/5/1993	DW-2-93	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
17	MW-2	6/12/1994	DW-2-94	Na-Ca-Cl-SO	silty-sand	19	<input checked="" type="checkbox"/>
18	MW-2	7/21/1995	DW-2-95	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
19	MW-2	7/24/1996	DW-2-96	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
20	MW-2	6/1/1997	DW-2-97	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
21	MW-2	7/24/1998	DW-2-98	Na-Ca-Cl-SO	silty-sand	19	<input type="checkbox"/>
8	MW-3	8/8/1992	MW-3-92	Ca-SO4-HCl	sandy-silt	20	<input type="checkbox"/>
9	MW-3	6/8/1993	MW-3-93	Ca-HCO3-SO	sandy-silt	20	<input type="checkbox"/>
10	MW-3	6/15/1994	MW-3-94	Ca-SO4-HCl	sandy-silt	20	<input type="checkbox"/>
11	MW-3	7/25/1995	MW-3-95	Ca-HCO3-SO	sandy-silt	20	<input type="checkbox"/>
12	MW-3	8/2/1996	MW-3-96	Ca-HCO3-SO	sandy-silt	20	<input type="checkbox"/>
13	MW-3	6/6/1997	MW-3-97	Ca-HCO3-SO	sandy-silt	20	<input checked="" type="checkbox"/>
22	MW-4	7/15/1992	DW-4-92	Ca-SO4	sandy-grave	21	<input type="checkbox"/>
23	MW-4	5/25/1993	DW-4-93	Ca-SO4	sandy-grave	21	<input type="checkbox"/>
24	MW-4	5/23/1994	DW-4-94	Ca-SO4	sandy-grave	21	<input type="checkbox"/>
25	MW-4	6/12/1995	DW-4-95	Ca-SO4	sandy-grave	21	<input checked="" type="checkbox"/>
26	MW-4	7/25/1996	DW-4-96	Ca-SO4	sandy-grave	21	<input type="checkbox"/>
27	MW-4	5/15/1997	DW-4-97	Ca-SO4	sandy-grave	21	<input type="checkbox"/>

The first column in these windows will always contain an ID value; each sample and station in your database will have a unique database ID value. This allows AquaChem to manage the data and perform internal calculations.

NOTE: The internal database ID value cannot be edited, nor can this column be removed from the active list. This ID value is automatically created when you create a new sample or station.

In addition to the ID column, there will be columns containing sample or station description parameters. These columns can be modified and the sorting options can be modified as well. For more details on sorting the active list, please see the **View > Options - Active List** section in Chapter 3.

The bottom of the **Active Sample/Stations** window contains the following three buttons:



The **[Sort]** button will load the sort options for the active list. This will allow you to change which parameters appear in the active list and their order.



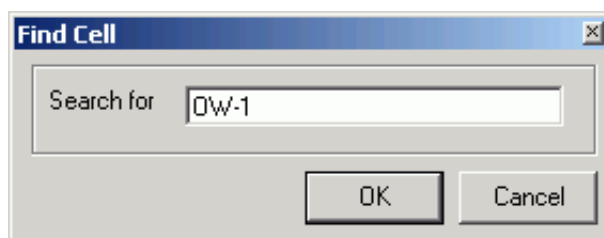
The **[New]** button will create a new sample or station, depending on which mode is active (i.e. which tab is selected).



The **[Delete]** button will delete the selected sample or station.

In order to edit the data for a specific sample or station, you need to open the **Sample Details** or **Station Details** window. These windows are explained in greater detail in the following sections.

The following shortcuts can be used to reach quickly a specific record without the more time consuming **Find** operation. Click in the column holding the field you want to search. Then press the first character of the searched expression on the keyboard. The program will jump to the first expression starting with the respective character. Alternatively, you may press Ctrl-F and type in the text string you want to search for. After pressing the **[OK]** button, the program will jump to the first matching record having a matching field.



If the **Find** function returned several records in a long list, these records may be scrolled through without hiding the unselected records. To do so, press Ctrl-S on your keyboard. This will initially take you to the first selected record, while pressing it again will scroll through all highlighted records sequentially.

1.5.2 Sample Details Window

The **Sample Details** window is a read/write window, which means data can be entered, saved, and read from this window. Individual samples can be viewed and edited using this window.

To load this window for one of the samples in your active list, you can:

- select a sample from the active list and **double-click** the left mouse button on it; OR
- select a sample from the active list and press the **<Enter>** key on your keyboard; OR
- select a sample from the active list and click **Sample > Edit** from the main menu; OR
- right-click the sample from the active list and select **[Edit]**.

An example of the **Sample Details** window is shown below:

To enter data in the **Sample Details** window, simply double-click in the desired field and type in the appropriate information. Alternatively, data can be imported into your database using the **Import** feature (see the **File > Import** section for more details).

The **Sample Details** window is separated into two frames: the top frame includes general details on the sample (**Sample** and **Station** tabs), and the bottom frame contains the **Measured**, **Calculated**, **Modeled**, and **Description** tabs.

Data can be entered for the **Sample** tab at the top of this window, and in the **Measured** and **Description** tabs in the bottom half of this window. Under the **Measured** Parameters tab, you will see the label **Parameter Group** with a corresponding combo box. This allows you to select different groups of Measured Parameters, and focus on just desired groups (for example you may want to view just Anions or Cations). The **Show analyzed values only** group will hide all parameters for which there is no data recorded, and display only those samples which have measured values. Parameter groups can be created and edited in the **File > Database** screen using the parameter group tab dialogue.

Parameter	Unit	Value	meq
Depth		5.0	
Temperature	°C	14.9	
pH (field)		7.15	
pH (lab)			
Eh	mV		
DO	mg/l	9.2	
TDC	mg/l		
TSS	mg/l		
El Cond.	uS/cm	1200	
TDS	mg/l	726	
Density	g/cm3		
Ca	mg/l	125	6.25
Mg	mg/l	22	1.1
Na	mg/l	80	3.45
K	mg/l	1.5	0.015
Cl	mg/l	125	3.45

For Sample, Station or Measured Parameter you may also right-mouse click on a grid cell in order to view the Parameter Details. The **Parameter Details** window displays all the meta data available for the selected parameter including description, formula weight, and the CAS Registry number.

The **Calculated** tab contains function values based on measured data from the current sample. These entries cannot be edited (this data is read-only). However you may define which of the available AquaChem functions should be displayed and what unit is to be displayed (e.g. for hardness) on this tab using the Sample Detail Options.

The data in the **Modeled** tab is obtained from PHREEQC simulations (as such, there will be no values for Modeled Parameters when you enter or import data into your database). There are two ways in which you can copy PHREEQC results into the **Modeled** tab:

- [1] Click the **PHREEQC** button at the bottom of the window, and PHREEQC will calculate the Saturation Indices for the available Modeled Parameters in the database. This will be done only for the current sample;
- [2] Select multiple samples in the Active Samples list, and use the menu option **Tools > Modeling > Calculate Sat. Indices and Activities**.

The scroll buttons at the bottom of the **Sample Details Window** can be used to scroll through the Sample Details for other samples:



The order of these buttons (from left to right) is as follows:

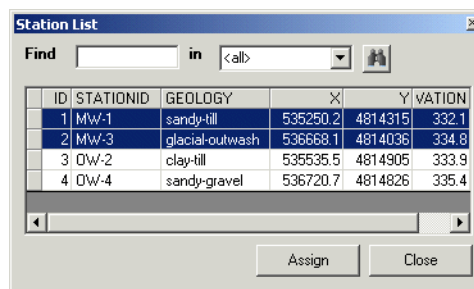
First sample - loads the sample details for the first sample in your active list.

Previous sample - loads the sample details for the previous sample in your active list.

Next sample - loads the sample details for the next sample in your active list.

Last sample - loads the sample details for the last sample in your active list.

The first field in the Sample Details window is the **Station ID**. As mentioned earlier, every sample must have a station assigned to it. To assign a station to a sample, click once in this field then click the **...** button which will appear near the right side of this field. Alternatively, you may click **Samples > Assign Station** from the main menu. This will load a list of available stations, similar to the dialogue shown to the right side.



From this dialogue, you may select a station directly from the list; or if you have a long list of stations, the **Find** feature at the top of this window can be helpful. Simply enter the Station ID or any other parameter from the station you are looking for into the **Find** field, and press the **Find** icon to run a search for this expression. If this expression might be found in several fields of the station table then you might want to choose a category from the combo box beside this field in order to narrow down the fields which are searched by the query.

Once you have located the desired station for this sample, press the **[Assign]** button at the bottom of this dialogue and this will return you to the Sample Details window.

When you are finished in the Sample Details window, press the **[Save]** button at the bottom to save new data and/or changes to your database. Once you are finished, press **[Close]** to return to the Active List.

The data under the **Station** tab is read-only, and as such cannot be edited. The Station tab contains information on the station which corresponds to the current sample. To edit the station parameters, open the **Station Details Window** as described in the next section.

Note that in the **Sample Details** section, fields with a defined length of greater than 100 characters will appear as a multiline field to make long texts, such as comments, readable. Also, the text fields in the **Sample Details** section may contain hyperlinks or file names. Double clicking such a field will open the respective file or link. It may be useful to link the original laboratory data files or PDF files from which the data was

extracted to respective samples. Even if the information contained in this file is not actually included in the AquaChem database, is then easily accessible.

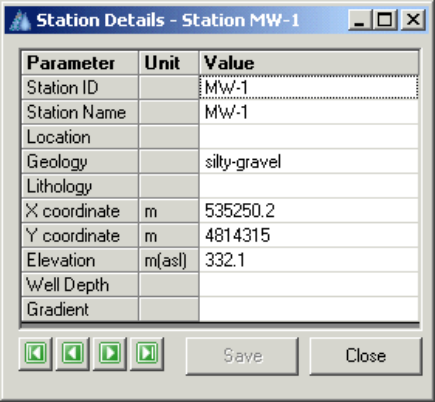
1.5.3 Station Details Window

The **Station Details** window is a read/write window, which means data can be entered, saved, and read from this window. Individual stations can be created, edited, or viewed using this window.

To load this window for one of the stations in your active list, you can:

- select the station from the active list, then **double-click** the left mouse button on it; OR
- select the station from the active list, then press the **<Enter>** key on your keyboard; OR
- select the station from the active list and click **Station > Edit** from the main menu; OR
- right-click on the station from the active list and select **[Edit]**.

An example of the **Station Details** window is shown below.



Parameter	Unit	Value
Station ID		MW-1
Station Name		MW-1
Location		
Geology		silty-gravel
Lithology		
X coordinate	m	535250.2
Y coordinate	m	4814315
Elevation	m(asl)	332.1
Well Depth		
Gradient		

At the bottom of the window, there are four small square buttons with arrows (left, right, up, down) and two larger buttons labeled 'Save' and 'Close'.

To enter data in the **Station Details** window, simply double-click in the desired field and type in the appropriate information. Alternatively, data can be imported into your database using the Import feature (see the **File > Import** section for more details). To save new data and/or changes to the database for this station, press the **[Save]** button at the bottom of this window. Once you are finished, press **[Close]** to return to the active list.

The scroll buttons at the bottom of this window are similar to the Sample Details window; these buttons can be used to scroll through the details for other stations in your active list.

Note that in the sample detail section, fields with a defined length of greater 100 will appear as a multiline field to make long texts such as comments readable. Also, the text fields in the sample detail section may contain hyperlinks or file names. When double clicking such a field, the respective file or link will be opened. Useful information to link to a station record may include an image of the station or a geological log.

1.5.4 Plots Window

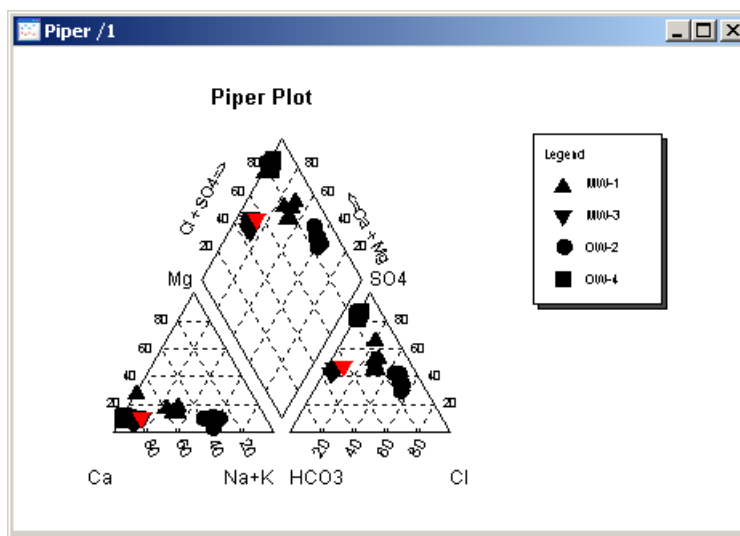
AquaChem provides a comprehensive selection of plotting techniques commonly used for aqueous geochemical data analysis and interpretation. Each of these plot types can be used to graphically represent information for all samples in the Active Samples List, or for selected samples only.

To create a new plot:

- Ensure the **Samples** tab is the current active window.
- Select **Plots > New** from the main menu.
- Choose the desired plot type from the list in this menu.
- Modify the plot options or click **[OK]** to accept the defaults.

This will create a Plot window displaying the selected plot for all or selected samples in the Active Samples List.

An example below shows a plot window containing a **Piper** plot:



Any samples selected in the **Active List** will be highlighted on the **Piper** plot. Shapes and sizes of the symbols can be modified and the plot options can be adjusted to show just the selected samples, or all the current active samples available in your database. In certain plots the data points may be labeled.

It is important to remember that the data plotted on all open plots are directly linked to the database samples. Any changes to the data are immediately reflected in each of the open graphs. Clicking a data point on the graph will highlight the corresponding sample in the Active samples list window (the corresponding data point in all other open plot windows will also be highlighted). This can be effective for identifying outlier points on the plot. Similarly, selecting a sample in the active list will highlight the corresponding data point on all open graphs. Changing the number of samples in the active list automatically updates ALL open plots.

For more details on the various Plots and their respective options, please refer to Chapter 4: Plots.

1.5.5 Table View

The **Table View** window is loaded when you select **View > Table View** from the main menu. You can then load any of the previously created table views, or use the **Create** option to design a new Table (spreadsheet) View.

For more details on the Table View options, please see the **View > Table View** section in Chapter 3.

1.5.6 Reports Window

A **Report** window provides reported and/or calculated information for a selected sample, group of samples, or all active samples in the database. The reports can be produced by selecting a sample from the active list and then selecting one of the report types from the **Reports Menu** option.

The text reports can be edited, printed, or saved to a .TXT, .CSV or .XLS file. AquaChem generates several types of reports. Using the **Report Designer**, you can create and customize your own reports, to display whatever data and/or calculations you desire.

For more details, please refer to Chapter 5: Reports.

1.5.7 Tools

AquaChem provides you with the following pre-defined data analysis tools:

- AquaChem Function
- Decay Calculator
- Find Missing Major Ion
- Formula Weight Calculator
- Volume Concentration Converter
- Special Conversions
- Species Converter
- Unit Conversions

- Calculate facies
- Corrosion and Scaling
- Oxygen solubility

There are also **QA/QC** checks, **Look Up Tables**, and options for the linking to the PHREEQC interface available under the **Tools Menu**. As well, there is a feature that allows you to create an input file for PHT3D modeling engine using the data entered in the database.

For more details, please refer to Chapter 6: Tools.

1.5.8 PHREEQC Interface

AquaChem includes a direct link to the USGS modeling program PHREEQC (version 2.12). You may also run the USGS graphical user interfaces (PHREEQCI or PHREEQC for Windows), giving access to more advanced options which are not available through the AquaChem interface.

For more details on PHREEQC and modelling, please refer to Chapter 7.

1.6 AquaChem Toolbar

This section describes each of the items in the AquaChem toolbar. Most toolbar buttons are context sensitive and react according to the active AquaChem window or dialogue. If there are no options available for the selected window or dialogue, the toolbar icons may become grey and inactive. The AquaChem toolbar is shown below.



For a short description of each item in the toolbar, place your mouse pointer over an icon and a hint will pop-up. The function of each toolbar item is described below:



New button creates a new database (only available if no other database is open)



Open button opens a database (only available if no other database is open)



Save button saves the current database file



Print button prints a plot, table, or a report



Copy button copies currently selected data, or copies a plot to the Windows Clipboard



Cut button cuts currently selected data



Paste button pastes currently copied (or cut) data



Edit button edits selected sample/station



Create button creates new sample/station



Delete button deletes selected sample/station



Find button finds samples/stations



Options button loads the options for sample/station list, Table View options, Report, or Plot window



Show all button shows all samples/stations in the active list



Omit all button hides all samples/stations in the active list



Show only selected button hides all samples/stations in the list that have not been selected



Omit selected button hides all selected samples/stations in the active list



Zoom out/Zoom in buttons are used to change the zoom extent in the **Map** and other X,Y plots



Identify button identifies sample data used on the selected plot(s)

2

Getting Started

This chapter is designed to serve as a ‘quick-start’ reference guide for those interested in getting started quickly using the most common features of AquaChem. You will follow a sample set of instructions, using the same data set that was used for the Demo database.

To begin, this chapter has been divided into sections for your convenience - feel free to read through the entire chapter, or jump directly to a section of interest.

Creating a New Database

- Importing Data
- Assigning Symbols

Creating Plots

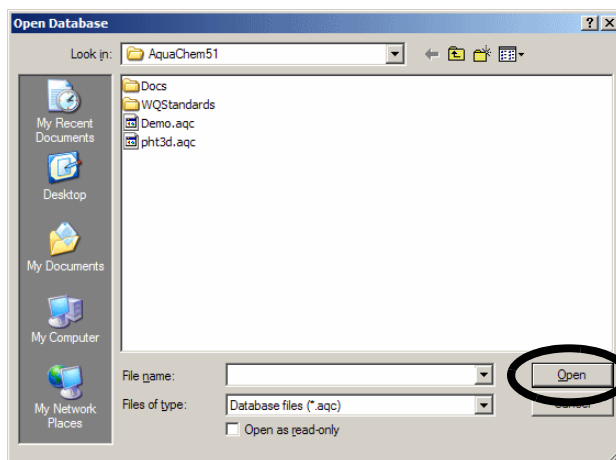
- Plot Options
- Printing Plots
- Exporting Plots as Graphics File

Creating Reports

- Saving Reports
- Printing Reports

2.1 Creating a New Database

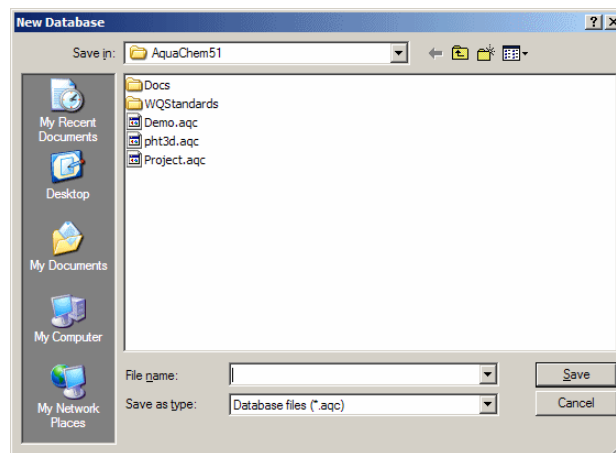
When you first load AquaChem, an **Open Database** dialogue will appear. You have the option of loading the Demo_Basic.AQC project, or to create a new project. To create a new project, press **[Cancel]** when this dialogue appears.



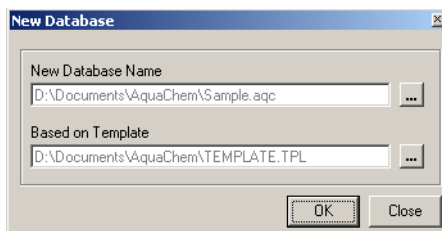
After pressing **[Cancel]**, you will see a blank AquaChem window indicating there is no database loaded.

To create a new database:

- Select **File** from the main menu and then select **New**. A **New Database** dialogue will appear as shown below.



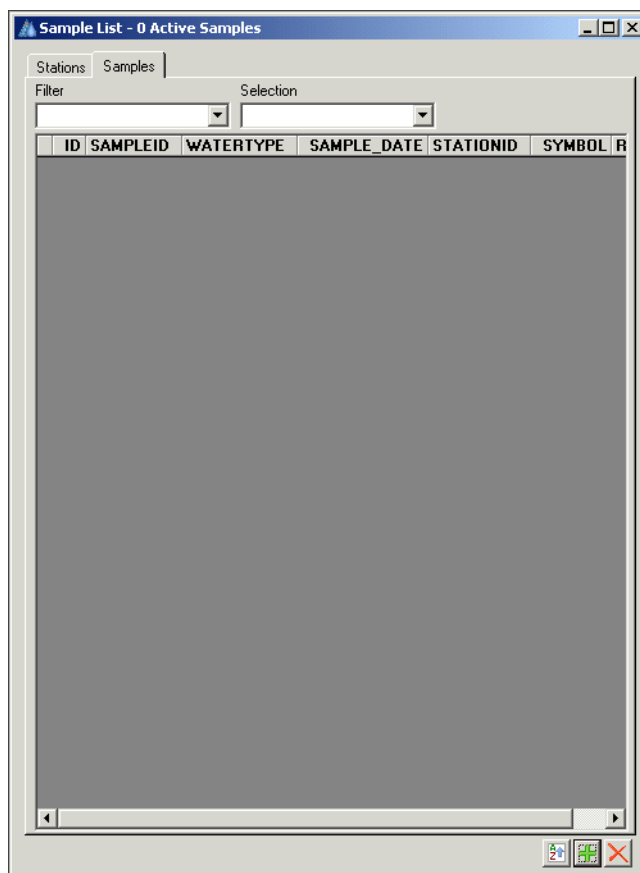
- Browse to the appropriate folder
- Type in the name of the new database, and click **[Save]**. (For this demonstration, the file will be called **Sample.aqc**).
- You will then see the following **New Database** confirmation dialogue.



- Below the **New Database Name** field, in the **Based on Template** field, you must specify a database template file to use for the creation of your database. If you are a new user to AquaChem, it is recommended that you use the **Template_Basic.TPL** file as the starting template for your database (the default). After becoming more experienced with the program, you can create new templates and use these for future databases.
- Click **[OK]** once you are finished entering the database (.AQC) and template (.TPL) name.

NOTE: It is recommended that database filenames contain a combination of letters and numbers only. The filename and directory path should NOT contain spaces, accents, or other special characters.

AquaChem will then automatically create a blank database file using the associated database template file containing the data structure for each record in the database. A blank database is shown below:



At this point you can begin to enter your sample data into the database. You have two options for data entry:



- Data can be entered manually; or
- Data can be imported from a text file.

The most convenient option for large data sets is to import the data from a text file; this will be demonstrated below. If you enter your data manually, be sure to first create a minimum of one station, so that you have a pre-existing station to assign to your samples. If the data is imported, the station data can be imported the same time as the sample date or separately.

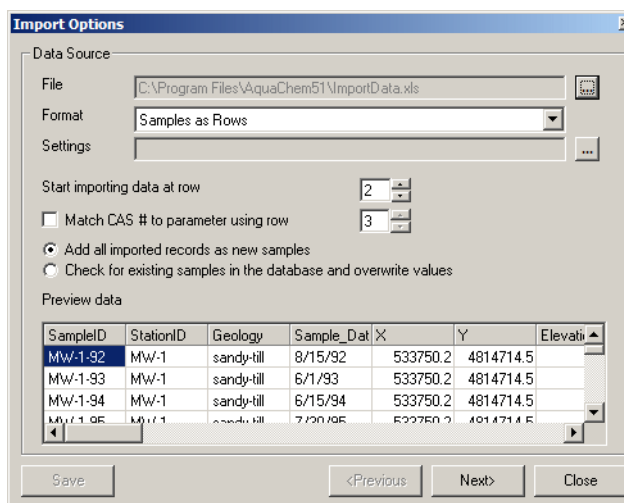
Before proceeding with the import option, you should ensure that your source file containing your water analysis data is properly formatted. For your convenience, an Excel template is provided with AquaChem, which includes the most common **Sample** and **Station** parameters. This file is called **template_2010.XLS** and can be found in the root AquaChem installation folder. If you wish, you may use this file for your data set. Simply enter your data, then **Save** the Excel file under a new name and then import your data to an AquaChem database.

2.1.1 Importing Data

To import sample data into your AquaChem database, please follow these steps:

- ☞ **File > Import > Samples/Stations** from the main menu
- ☞  button beside the **File** field to locate the source file
- ☞ **ImportData.xls** (located in the *Documents\AquaChem* directory).
- ☞ **[Open]**
- ☞  to select the import **Format** for the source file; **Samples as Rows** or **Samples as Columns**. For this demonstration, the **Samples as Rows** mode will be used.

You should now see the ImportData.xls file name in the **File** field, and a preview of file in the lower half of the dialogue (as shown below):



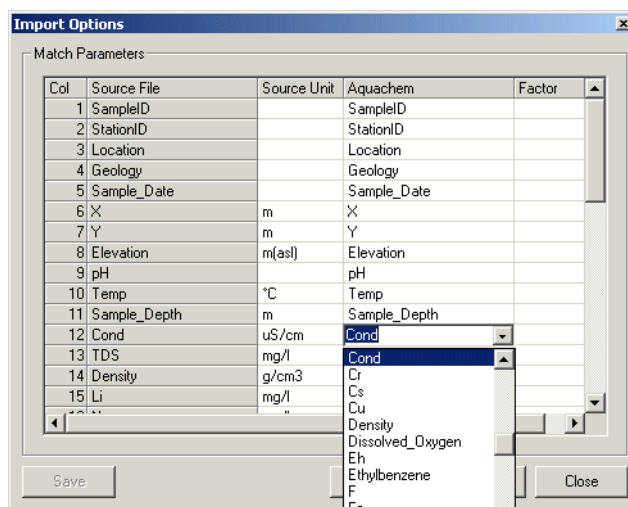
SampleID	StationID	Geology	Sample_Dat	X	Y	Elevation
MW-1-92	MW-1	sandy-till	8/15/92	533750.2	4814714.5	
MW-1-93	MW-1	sandy-till	6/1/93	533750.2	4814714.5	
MW-1-94	MW-1	sandy-till	6/15/94	533750.2	4814714.5	
MW-1-95	MW-1	sandy-till	7/20/95	533750.2	4814714.5	

The remaining import options can be left as is.

- ☞ **[Next>]**

In the next dialogue, AquaChem requires you to match the fields in your file to the required AquaChem parameter fields. For the **ImportData.xls** file, the column headings have already been prepared such that they precisely match up with the required AquaChem field headings. Therefore, there is no need to do manual matching. However, if there are blank fields in the AquaChem column, then this indicates that the parameter names in the file are not identical to the AquaChem parameter names, and

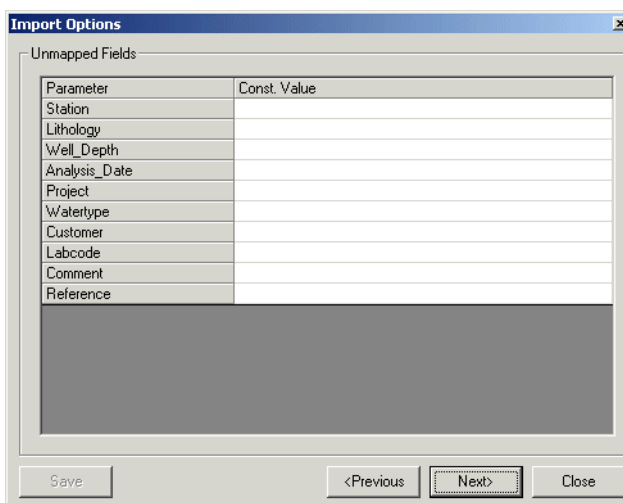
the fields could not be identified. For example, if your file contained the parameter name '**Conductivity**', you must match this up to the AquaChem parameter Internal Key '**Cond**'.



If necessary, please refer to the Import section in Chapter 3 for more details on matching parameters during the import.

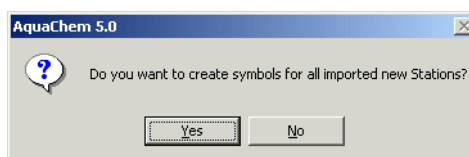
☞ **[Next>]**

The next dialogue contains a list of the AquaChem parameters which were not matched up to parameters in the source file. This dialogue allows you to enter constant values for these parameters (optional). For example, if all of the imported samples belong to the same project or the same location and this information is not included in the source data file, you may fill the respective fields during this step of the import. To skip this step, leave these fields blank.



☞ **[Next>]**

You will then see the following warning prompt:



This option allows you to automatically create one symbol for each unique station in the imported source file. These symbols will be created in the Station symbol group. This is a very helpful feature, which can aid in plotting your sample data later on.

☞ **[No]** to pass on this feature.

The last dialogue in the import options, contains a summary of the import routine and gives you the opportunity to use the **[Save]** button to save the import settings. Please check that all of your samples, stations, and parameters have been successfully imported.

☞ **[Close]**

After successfully importing the data, you will return to the main database window showing your active **Samples** and **Stations**. For this data set, there should be 4 stations, with 7 samples taken from each station location, for a total of 28 samples.



2.1.2 Assigning Symbols

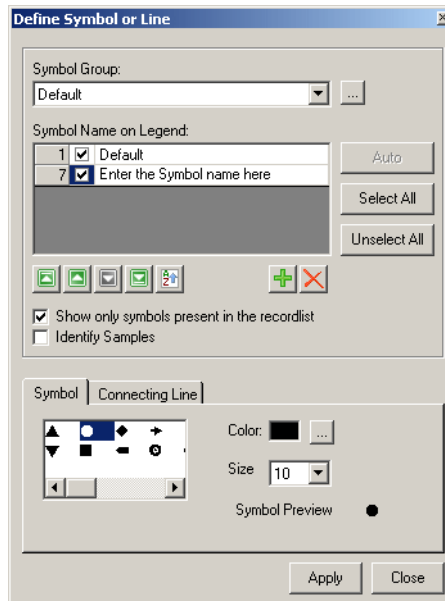
The next step to building a new AquaChem project is to assign symbols to your samples.

By default, a new database will include two symbol groups: a **Default** symbol group with just one symbol assigned to each sample, and a **Station** symbol group, with a unique symbol for each unique station in the database; the station symbols are automatically assigned to the appropriate corresponding samples. You may create new symbol groups, or add symbols to one of the existing symbol groups.

For this demonstration, you will add symbols to the **Default** group.

Follow the instructions below to create new symbols.

- ☞ **Plots** from the main menu and then select **Define Symbol or Line**
- ☞  and ensure the **Default** symbol group is selected.
- ☞  button and a new field will be added to the list of symbols



type: **MW-1** in the new line that is added

- ☞ **<Enter>** (on your keyboard) to accept the new name

You will now add three more symbols:

- ☞  button

type: **MW-3** (for the new symbol)

- ☞ **<Enter>** on your keyboard

☞  button

type: **OW-2** (for the new symbol)

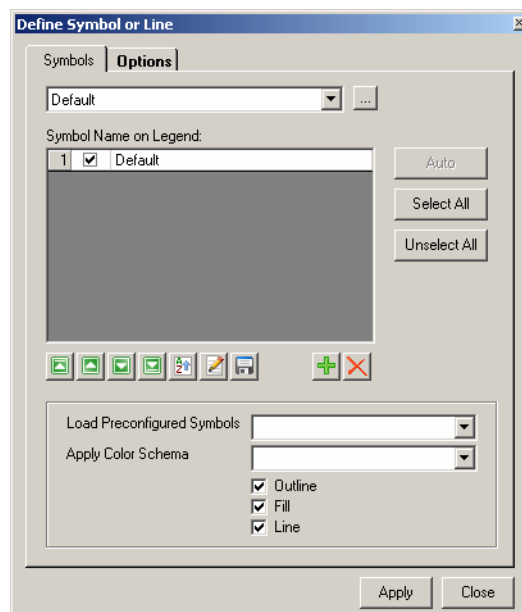
☞ **<Enter>**

☞  button

type: **OW-4** (for the new symbol)

☞ **<Enter>**

Once you are finished, you should have five symbols listed for the **Default** symbol group, as seen in the figure below.



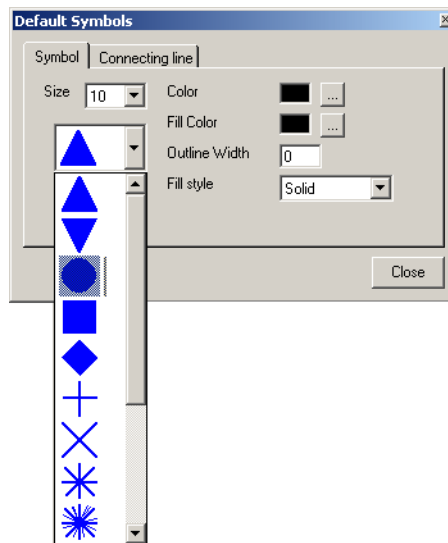
Next you need to define the necessary symbol properties:


☞ **Symbol #2 (MW-1)** from the list

☞  **Edit** button

The **Default Symbols** dialog will appear on your screen.

- ☞ Choose a symbol character from the list of available symbols . Simply click on the desired new symbol shape



- ☞ Choose a **Color** for the new symbol. To access the color options press the  button beside **Color**, and select a color from the list of available colors
- ☞ Choose a **Size** for the symbol; the font sizes are available from the combo box beside **Size**. (Recommended size is 12pt.)
- ☞ **[Close]** button, once you are finished.

Repeat these steps for the three other new symbols that were created.

Once you are finished creating the symbols and defining the symbol properties,

- ☞ **[Apply]**

- ☞ **[Close]**

In your active samples list, you are ready to assign these symbols to the samples.

ID	SAMPLEID	WATERTYPE	SAMPLE_DATE	STATIONID	SYMBOL	REP
1	MW-1-92	Ca-Na-SO4-Cl	8/15/1992	MW-1	1	<input checked="" type="checkbox"/>
2	MW-1-93	Ca-Na-SO4-CHCO3	6/1/1993	MW-1	1	<input checked="" type="checkbox"/>
3	MW-1-94	Ca-Na-SO4-Cl	6/15/1994	MW-1	1	<input checked="" type="checkbox"/>
4	MW-1-95	Ca-Na-SO4-CHCO3	7/30/1995	MW-1	1	<input checked="" type="checkbox"/>
5	MW-1-96	Ca-Na-SO4-CHCO3	7/28/1996	MW-1	1	<input checked="" type="checkbox"/>
6	MW-1-97	Ca-Na-SO4-CHCO3	6/15/1997	MW-1	1	<input checked="" type="checkbox"/>
7	MW-1-98	Ca-Na-SO4-CHCO3	8/1/1998	MW-1	1	<input checked="" type="checkbox"/>
8	MW-3-92	Ca-SO4-HCO3	8/8/1992	MW-3	1	<input type="checkbox"/>
9	MW-3-93	Ca-HCO3-SO4	6/8/1993	MW-3	1	<input type="checkbox"/>
10	MW-3-94	Ca-SO4-HCO3	6/15/1994	MW-3	1	<input type="checkbox"/>
11	MW-3-95	Ca-HCO3-SO4	7/25/1995	MW-3	1	<input type="checkbox"/>
12	MW-3-96	Ca-HCO3-SO4	8/2/1996	MW-3	1	<input type="checkbox"/>
13	MW-3-97	Ca-HCO3-SO4	6/6/1997	MW-3	1	<input type="checkbox"/>
14	MW-3-98	Ca-HCO3-SO4	7/30/1998	MW-3	1	<input type="checkbox"/>
15	OW-2-92	Na-Ca-Cl-SO4	8/1/1992	OW-2	1	<input type="checkbox"/>
16	OW-2-93	Na-Ca-Cl-SO4	6/5/1993	OW-2	1	<input type="checkbox"/>
17	OW-2-94	Na-Ca-Cl-SO4	6/12/1994	OW-2	1	<input type="checkbox"/>
18	OW-2-95	Na-Ca-Cl-SO4	7/21/1995	OW-2	1	<input type="checkbox"/>
19	OW-2-96	Na-Ca-Cl-SO4	7/24/1996	OW-2	1	<input type="checkbox"/>
20	OW-2-97	Na-Ca-Cl-SO4	6/1/1997	OW-2	1	<input type="checkbox"/>
21	OW-2-98	Na-Ca-Cl-SO4	7/24/1998	OW-2	1	<input type="checkbox"/>
22	OW-4-92	Ca-SO4	7/15/1992	OW-4	1	<input type="checkbox"/>
23	OW-4-93	Ca-SO4	5/25/1993	OW-4	1	<input type="checkbox"/>
24	OW-4-94	Ca-SO4	5/23/1994	OW-4	1	<input type="checkbox"/>
25	OW-4-95	Ca-SO4	6/12/1995	OW-4	1	<input type="checkbox"/>
26	OW-4-96	Ca-SO4	7/25/1996	OW-4	1	<input type="checkbox"/>
27	OW-4-97	Ca-SO4	5/15/1997	OW-4	1	<input type="checkbox"/>
28	OW-4-98	Ca-SO4	7/12/1998	OW-4	1	<input type="checkbox"/>

- ☞ Select the first seven samples in your sample list. To do so, click on the first sample, hold down your left mouse button, and drag your mouse down to the 7th sample.
- ☞ **Samples** from the main menu and then select **Assign Symbol** from the main menu (or use the Assign Symbol icon in the tool bar).

The **Assign Symbol** dialogue will list the available Symbol group names and the corresponding symbol, as defined earlier.

- ☞ Symbol **02 MW-1**
- ☞ **[Assign]**

Symbol Name	Symbol Preview
01 Default	▲
02 MW-1	■
03 MW-3	◆
04 OW-2	●
05 OW-4	▲

You will now assign symbols to the next group of samples.

- ☞ Without closing the **Assign Symbol** dialogue, activate the Sample list and select the samples from the **MW-3** Station- samples 8 to 14. The **Assign Symbol** dialogue will remain on top; move it if it is blocking the samples.
- ☞ Click in the **Assign Symbol** box to make it active.
- ☞ Symbol **03 MW-3**
- ☞ **[Assign]**

Repeat these steps to assign symbols to the remaining samples for stations **OW-2** and **OW-4**. Close the **Assign Symbol** dialogue by clicking the **[Close]** button when you are done.

Once you are finished, each of your samples should have a corresponding symbol which relates to the location of that sample. In the active list of samples, there is a column heading **Symbol** which indicates the Symbol group that is assigned to each sample. You should see Symbols 2, 3, 4, and 5 in your sample list, as shown below.

ID	SAMPLEID	WATERTYPE	SAMPLE_DATE	STATIONID	SYMBOL	EP
1	MW-1-92	Ca-Na-SO4-Cl	8/15/1992	Mw-1	2	<input type="checkbox"/>
2	MW-1-93	Ca-Na-SO4-CHCO3	6/1/1993	Mw-1	2	<input type="checkbox"/>
3	MW-1-94	Ca-Na-SO4-Cl	6/15/1994	Mw-1	2	<input type="checkbox"/>
4	MW-1-95	Ca-Na-SO4-CHCO3	7/30/1995	Mw-1	2	<input type="checkbox"/>
5	MW-1-96	Ca-Na-SO4-CHCO3	7/28/1996	Mw-1	2	<input type="checkbox"/>
6	MW-1-97	Ca-Na-SO4-CHCO3	6/15/1997	Mw-1	2	<input type="checkbox"/>
7	MW-1-98	Ca-Na-SO4-CHCO3	8/1/1998	Mw-1	2	<input type="checkbox"/>
8	MW-3-92	Ca-SO4-HCO3	8/8/1992	Mw-3	3	<input type="checkbox"/>
9	MW-3-93	Ca-HCO3-SO4	6/8/1993	Mw-3	3	<input type="checkbox"/>
10	MW-3-94	Ca-SO4-HCO3	6/15/1994	Mw-3	3	<input type="checkbox"/>
11	MW-3-95	Ca-HCO3-SO4	7/25/1995	Mw-3	3	<input type="checkbox"/>
12	MW-3-96	Ca-HCO3-SO4	8/2/1996	Mw-3	3	<input type="checkbox"/>
13	MW-3-97	Ca-HCO3-SO4	6/6/1997	Mw-3	3	<input type="checkbox"/>
14	MW-3-98	Ca-HCO3-SO4	7/30/1998	Mw-3	3	<input type="checkbox"/>
15	OW-2-92	Na-Ca-Cl-SO4	8/1/1992	OW-2	4	<input type="checkbox"/>
16	OW-2-93	Na-Ca-Cl-SO4	6/5/1993	OW-2	4	<input type="checkbox"/>
17	OW-2-94	Na-Ca-Cl-SO4	6/12/1994	OW-2	4	<input type="checkbox"/>
18	OW-2-95	Na-Ca-Cl-SO4	7/21/1995	OW-2	4	<input type="checkbox"/>
19	OW-2-96	Na-Ca-Cl-SO4	7/24/1996	OW-2	4	<input type="checkbox"/>
20	OW-2-97	Na-Ca-Cl-SO4	6/1/1997	OW-2	4	<input type="checkbox"/>
21	OW-2-98	Na-Ca-Cl-SO4	7/24/1998	OW-2	4	<input type="checkbox"/>
22	OW-4-92	Ca-SO4	7/15/1992	OW-4	5	<input type="checkbox"/>
23	OW-4-93	Ca-SO4	5/25/1993	OW-4	5	<input type="checkbox"/>
24	OW-4-94	Ca-SO4	5/23/1994	OW-4	5	<input type="checkbox"/>
25	OW-4-95	Ca-SO4	6/12/1995	OW-4	5	<input type="checkbox"/>
26	OW-4-96	Ca-SO4	7/25/1996	OW-4	5	<input type="checkbox"/>
27	OW-4-97	Ca-SO4	5/15/1997	OW-4	5	<input type="checkbox"/>
28	OW-4-98	Ca-SO4	7/12/1998	OW-4	5	<input type="checkbox"/>

The samples are now ready to be plotted.

2.2 Creating Plots

AquaChem allows you to plot your sample data numerous different plot types. For this exercise, you will create a simple Piper plot which distinctly shows the four different sample groups. For more details on the AquaChem plots and the plot options, please refer to Chapter 4.

In the **Sample Lists**,

- ☞ Highlight all samples in the list, by selecting the first sample in the list and dragging the mouse down the list while holding the left-mouse button.

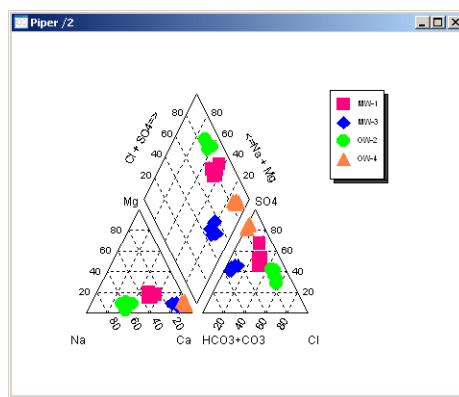
ID	Station	Date	Code	WATERTY	GEOLOGY	SYMBOL	REP
1	MW-1	8/15/1992	Mw-1-92	Ca-Na-SO4	sandy-til	2	■
2	MW-1	6/1/1993	Mw-1-93	Ca-Na-SO4	sandy-til	2	■
3	MW-1	6/15/1994	Mw-1-94	Ca-Na-SO4	sandy-til	2	■
4	MW-1	7/30/1995	Mw-1-95	Ca-Na-SO4	sandy-til	2	■
5	MW-1	7/28/1996	Mw-1-96	Ca-Na-SO4	sandy-til	2	■
6	MW-1	6/15/1997	Mw-1-97	Ca-Na-SO4	sandy-til	2	■
7	MW-1	8/1/1998	Mw-1-98	Ca-Na-SO4	sandy-til	2	■
8	MW-3	8/8/1992	Mw-3-92	Ca-SO4-HCl	glacial-outw	3	■
9	MW-3	6/8/1993	Mw-3-93	Ca-HCO3-Si	glacial-outw	3	■
10	MW-3	6/15/1994	Mw-3-94	Ca-SO4-HCl	glacial-outw	3	■
11	MW-3	7/25/1995	Mw-3-95	Ca-HCO3-Si	glacial-outw	3	■
12	MW-3	8/2/1996	Mw-3-96	Ca-HCO3-Si	glacial-outw	3	■
13	MW-3	6/6/1997	Mw-3-97	Ca-HCO3-Si	glacial-outw	3	■
14	MW-3	7/30/1998	Mw-3-98	Ca-HCO3-Si	glacial-outw	3	■
15	DW-2	8/1/1992	Dw-2-92	Na-Ca-Cl-SC	clay-til	4	■
16	DW-2	6/5/1993	Dw-2-93	Na-Ca-Cl-SC	clay-til	4	■
17	DW-2	6/12/1994	Dw-2-94	Na-Ca-Cl-SC	clay-til	4	■
18	DW-2	7/21/1995	Dw-2-95	Na-Ca-Cl-SC	clay-til	4	■
19	DW-2	7/24/1996	Dw-2-96	Na-Ca-Cl-SC	clay-til	4	■
20	DW-2	6/1/1997	Dw-2-97	Na-Ca-Cl-SC	clay-til	4	■
21	DW-2	7/24/1998	Dw-2-98	Na-Ca-Cl-SC	clay-til	4	■
22	DW-4	7/15/1992	Dw-4-92	Ca-SO4	sandy-grave	5	■
23	DW-4	5/25/1993	Dw-4-93	Ca-SO4	sandy-grave	5	■
24	DW-4	5/23/1994	Dw-4-94	Ca-SO4	sandy-grave	5	■
25	DW-4	6/12/1995	Dw-4-95	Ca-SO4	sandy-grave	5	■
26	DW-4	7/25/1996	Dw-4-96	Ca-SO4	sandy-grave	5	■
27	DW-4	5/15/1997	Dw-4-97	Ca-SO4	sandy-grave	5	■
28	DW-4	7/12/1998	Dw-4-98	Ca-SO4	sandy-grave	5	■

- ☞ **Plots** from the main menu, then select **New** and then **Piper**

A **Piper Plot Options** dialogue will appear with default plot settings. The following section describes some of the generic plot options.

To create the plot using the default plot options, click **[OK]**.

The plot should be created similar to the one shown below:



You can modify the plot options by right-clicking on the plot to load the Plot options dialogue.

If you are satisfied with your plot, proceed to the next section titled **Printing Plots**.

2.2.1 Plot Options

There are many common graphical features and options for each plot. When you select any of the graph types to plot (for example Piper plot), a plot options dialogue will appear with default settings for all of the necessary parameters and settings. For most plot types, the **Plot Options** dialogue contains links to the following dialogues: **Parameters**, **Title**, **Symbols**, **Label**, **Axis**, and **Legend** (not all dialogue options are present on the Piper plot options dialogue. For more information, see Chapter 4)

Cations			Anions		
Parameter	Label	Factor	Parameter	Label	Factor
Na	Na	1	HCO3+CO3	HCO3+	1
Ca	Ca	1	Cl	Cl	1
Mg	Mg	1	SO4	SO4	1

Plot	
Title	...
<input checked="" type="checkbox"/> Legend	...
Symbols	...
<input type="checkbox"/> Labels	...
<input checked="" type="checkbox"/> Labeled Ticks	Δ
Axis Titles	Δ
<input checked="" type="checkbox"/> Show Grid	


Buttons: Set Default, Apply, OK, Cancel

The main Plot Options dialogue contains information on the parameters/series used in the plot and provides the link buttons to access various dialogues described below, which are used to customize the plot.

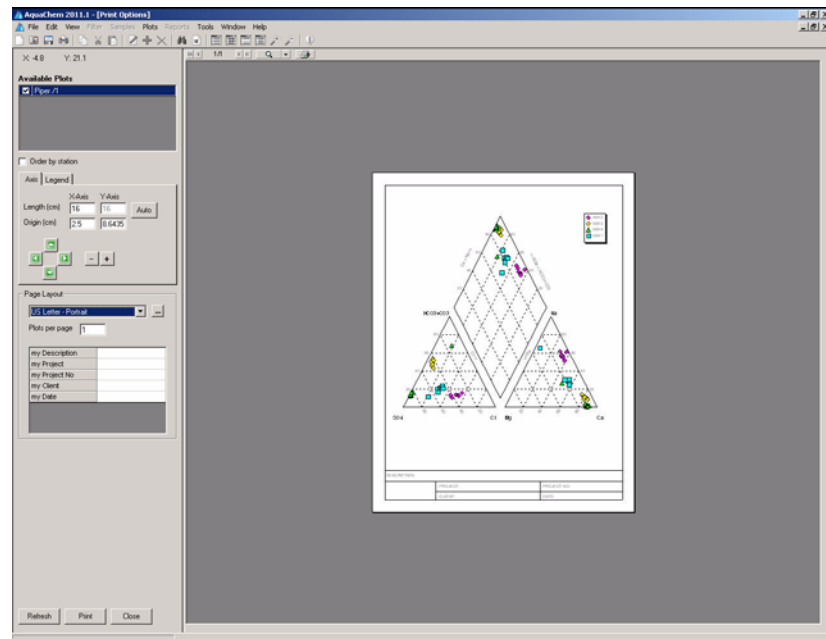
- **Title Dialogue** - contains options for **Plot Title** font size, **Position**, and **Alignment**.
- **Symbols Dialogue** - contains options for symbols used in the plot, **Visibility**, **Scaled Symbol Size** options, etc.
- **Legend Dialogue** - contains options for displaying a **Plot Legend**, Legend **Title**, and display features.
- **Labels Dialogue** - contains options for data point labeling and positioning.
- **Axis Dialogue** - contains options to customize the axes.
- **Line Dialogue** - contains options to add and customize lines on the plot.

2.2.2 Printing Plots

Once you have prepared the desired plot(s), you are ready to print.

☞ **File** from the main menu and select **Print**, or click the Print icon  in the tool bar.

You should then see a **Print Options** window, as seen below.

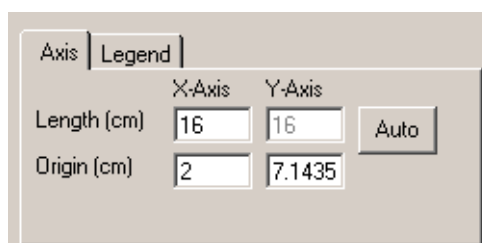


The Print Options allow you to choose which plots will be printed, their position, size, and which plot template will be used.

A list of **Available Plots** will appear in the upper-left corner of this window. This list represents the plots which are currently open in AquaChem. Select the plots you want to print by placing a check mark in the box beside the appropriate plot name. The piper plot is already selected by default.

If you have generated several plots you have the option to order them by station by select this check box under the list of available plots.

The plot will be automatically sized to fit the page layout. If the default page settings are not suitable, you can manually change the position of each plot using the options provided in the **Axis** tab (**Length** and **Origin**).





The screenshot shows the 'Axis' tab of a dialog box. It has two sub-tabs: 'X-Axis' and 'Y-Axis'. Under 'X-Axis', there are two input fields: 'Length (cm)' with the value '16' and 'Origin (cm)' with the value '2'. Under 'Y-Axis', there are two input fields: 'Length (cm)' with the value '16' and 'Origin (cm)' with the value '7.1435'. To the right of these fields is an 'Auto' button. The 'Legend' tab is also visible at the top of the dialog box.

If you want to display plots on several pages proceed as follows:

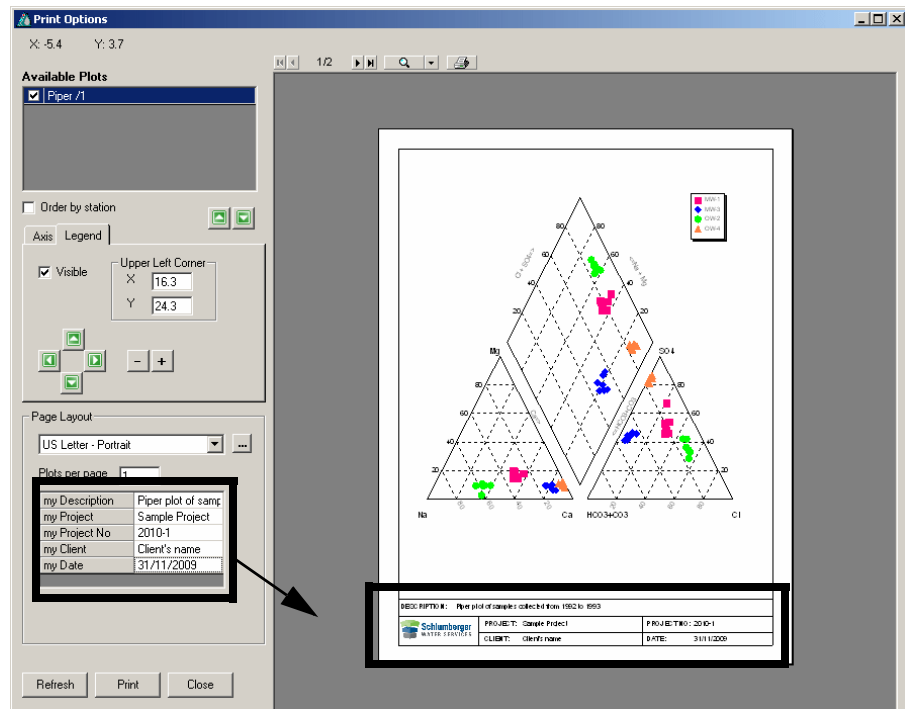
- Enter the number of plots per page.
- You will be asked to confirm the number of rows and columns on you page. For example, if you wish to display 8 plots and you enter 2 rows and 2 column, the plots will be distributed on 2 pages.

Next, you can select the page layout by choosing a print template. The print template will allow you to enter information on the page footer such as project information, sample information, and company information.

Under the **Page Layout** options (on the left side of the window), you will see the default template setting is **US-Letter-Portrait**.

  from the combo box to see all available templates.

You can see a list of plot descriptors in the **Page Layout** dialogue. They are linked to the fields on the report and you can see the changes made as soon as you leave the field you were typing in:



Next, you will fill in the project specific plot description fields under the **Page Layout** options. Press the <Enter> key after each entry:

DESCRIPTION:

type: Piper plot of samples collected from 1992 to 1998

PROJECT:

type: Sample Project

PROJ #:

type: 2010-1

CLIENT:

type: Client's name.

DATE:

type: Current date

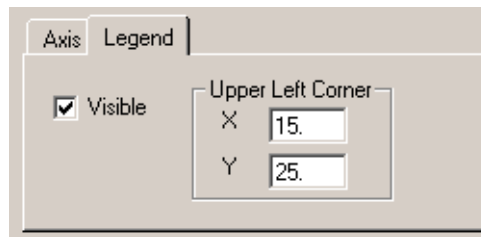
NOTE: The Schlumberger Water Services logo shown in the bottom of the page can be easily replaced with your own company logo. This is done using the **Template Designer** option, which can be accessed by clicking the **...** button beside the Template combo box. This option is not explored in this exercise. For more information, please refer to Template Designer section in Chapter 3.

Next, if you have added a legend to your plot, you can add this legend to the printed page, and position it on the page.

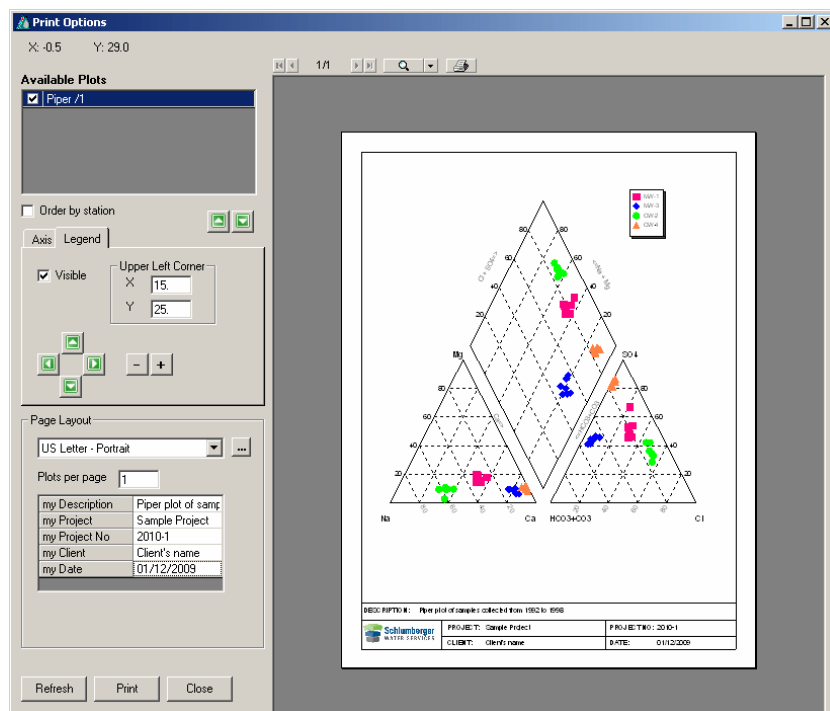
- ☞ **Legend** tab (below the list of **Available Plots**, and beside the **Axis** tab).
- ☞ **Visible** (click once in the check box) to activate the legend for the Piper Plot.

The **Piper** plot legend will appear in the upper-left corner of the page. To move the legend,

- ☞ **X-Axis** field and enter a value of 15.
- ☞ **Y-Axis** field and enter a value of 25.



If you have loaded the plot successfully, your display should be similar to the one shown below:



- ☞ **[Print]** button (in the lower left corner) to send the plot to a printer.
- ☞ **[Close]** to close the print options window and return to the AquaChem active samples list.

2.2.3 Exporting Plots as Graphics File

Upon returning to the main AquaChem window, you will see that the active sample list window is open, and you should still have the Piper Plot window open. In addition to printing the Piper plot, you will now save this plot as a graphics file so that it may be inserted into an external report.



- ☞ **Piper plot window** to make this the active window
- ☞ **File** from the main menu, and then choose **Export**, and finally, **Image**.

You will then be prompted to enter a filename for the plot. Browse to the desired folder on your computer, then enter a name for the file. For this demonstration, you will save the file in the AquaChem folder.

type: **Piper** as the filename for the plot

 **[Save]**

The plot will be saved as a .WMF (Windows MetaFile) graphics file. This file can be opened using most conventional graphics editors, or the file can be directly inserted into a word processor application.


  button in the upper-right corner of the **Piper Plot window**, to close this plot window.

In the next section, the Report options will be briefly demonstrated.

2.3 Creating Reports

AquaChem allows you to choose from 9 data analysis reports and perform 6 different statistical tests on your data. For this demonstration, you will briefly examine the generic Report options and create a Statistics Report. The **Summary Statistics** report provides a statistical summary of selected parameters for all active samples in your database. For more details on Reports, please refer to Chapter 5.

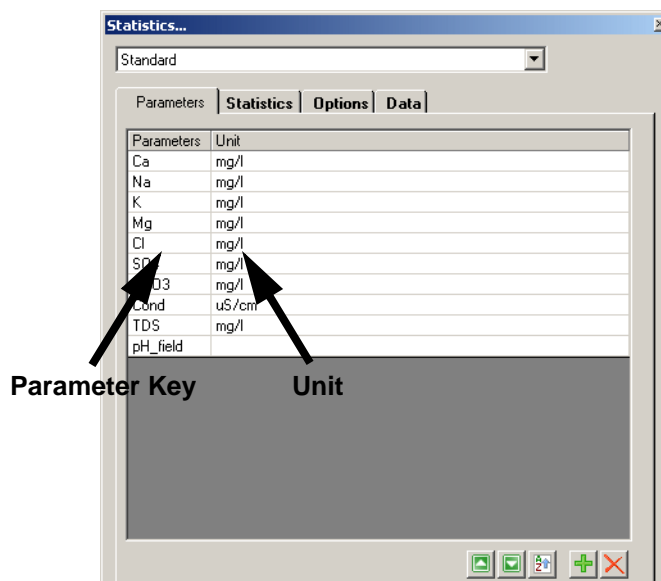
To create a **Summary Statistics** report for all samples in the active list:

 **Reports** from the main menu and then choose **Statistics**, and finally **Summary Statistics**


A dialogue will load that will allow you to customize your **Summary Statistics** report. This dialogue contains four tabs: **Parameters**, **Statistics**, **Options**, and **Data**.

Parameters tab

The first tab that loads will allow you to specify which parameters you wish to use in your **Summary Statistics** report.




For this example, you will add benzene, Sum of Anions and Sum of Cations to the list of parameters.

- ☞  button
- ☞ Select **Measured Values** from the combo box.
- ☞ Select **Benzene** from the list of parameters.
- ☞ **[Select]**
- ☞ Select **Calculated Values** from the combo box at the top of the dialogue.
- ☞ Select **Sum of Anions** from the list.
- ☞ **[Select]**
- ☞ Select **Sum of Cations** from the list.
- ☞ **[Select]**
- ☞ **[Close]**

If you wish to add more parameters, you can do so in the manner described above.

To remove a parameter from the list,

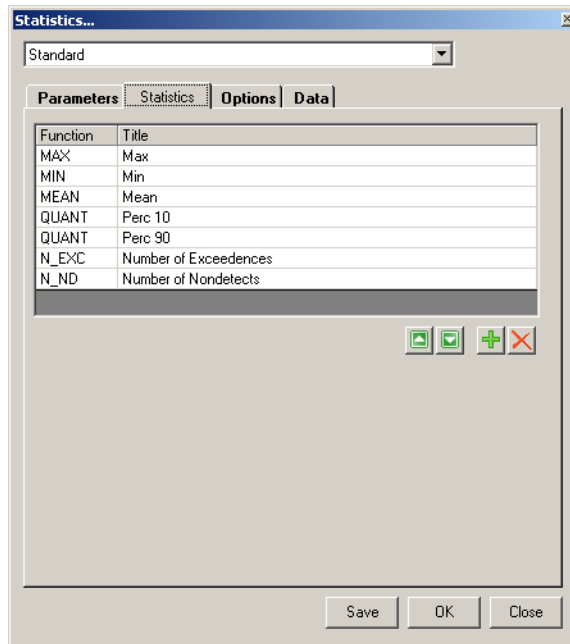
- ☞ Select the parameter to be removed
- ☞  button

Once all parameters are selected, you can chose what statistics to perform.

☞ **Statistics** tab.

Statistics tab

The **Statistics** tab allows you to select the summary statistics that will be performed on the data for the parameters you selected in the previous tab.



For this example, leave the statistics that are selected by default, however if you wish to add more statistics you can do so in the same manner as adding parameters. Selecting certain statistics will require you to enter additional information. For example, when selecting the Quantile statistic (QUANT), you will need to enter the percentage of the quantile.

When you are finished, move on to the **Options** tab.

Options tab

This tab allows you to select the output orientation (parameters as rows/statistics as columns or parameters as columns/statistics as rows) as well as providing you with the option to **Break by symbol**. If this option is selected, the summary report will be generated for every group of samples that bear the same symbol, i.e. every station in this example.

Statistics...

Standard

Parameters Statistics Options Data

Parameter as Rows

☐ Break by symbol

☒ Show guidelines

Show MDL from sample

Save OK Close

For this example, check the box beside **Break by symbol** and move on to the **Data** tab. This will create separate statistics for all samples assigned to the same symbol.

Data tab

The **Data** tab provides a preview of the data that will be used for the summary statistics report.

Statistics...

Standard

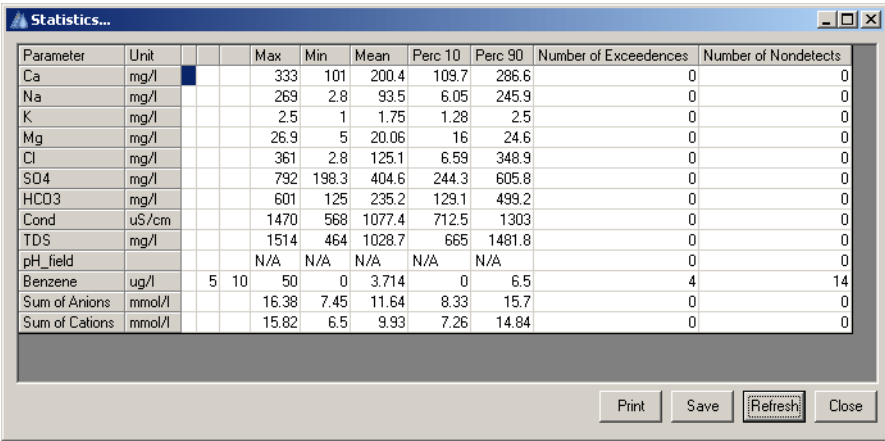
Parameters Statistics Options Data

Ca	Na	K	Mg	Cl	SO
mg/l	mg/l	mg/l	mg/l	mg/l	mg
125.0	80.0	1.5	22.0	125.0	550.
135.0	68.0	1.4	26.0	121.0	325.
135.0	86.0	2.5	23.4	128.0	325.
115.0	81.0	1.5	26.9	119.0	275.
105.0	76.0	2.0	26.2	115.0	254.
110.0	72.0	2.0	18.3	108.0	221.
101.0	75.0	2.5	17.5	111.0	216.
256.0	50.0	1.7	19.0	46.0	368.
268.0	35.0	1.5	20.4	42.0	346.
265.0	53.0	2.5	19.0	65.0	376.
275.0	39.0	2.0	12.5	30.0	348.
279.0	32.0	1.1	11.8	45.0	348.
286.0	45.0	1.3	18.2	32.0	351.
288.0	62.0	1.8	21.6	45.0	374.
115.0	225.0	1.3	5.0	286.0	198.
117.6	235.0	2.5	23.3	320.0	315.
126.0	245.0	1.0	20.7	351.0	308.
132.0	242.0	1.3	21.9	358.0	302.
162.0	248.0	2.3	24.0	348.0	418.

Save OK Close

☞ **OK** button.

The **Statistics** report window should be shown on your display, similar to the one below.



Parameter	Unit			Max	Min	Mean	Perc 10	Perc 90	Number of Exceedences	Number of Nondetects
Ca	mg/l			333	101	200.4	109.7	286.6	0	0
Na	mg/l			269	2.8	93.5	6.05	245.9	0	0
K	mg/l			2.5	1	1.75	1.28	2.5	0	0
Mg	mg/l			26.9	5	20.06	16	24.6	0	0
Cl	mg/l			361	2.8	125.1	6.59	348.9	0	0
SO4	mg/l			792	198.3	404.6	244.3	605.8	0	0
HCO3	mg/l			601	125	235.2	129.1	499.2	0	0
Cond	uS/cm			1470	568	1077.4	712.5	1303	0	0
TDS	mg/l			1514	464	1028.7	665	1481.8	0	0
pH_field				N/A	N/A	N/A	N/A	N/A	0	0
Benzene	ug/l	5	10	50	0	3.714	0	6.5	4	14
Sum of Anions	mmol/l			16.38	7.45	11.64	8.33	15.7	0	0
Sum of Cations	mmol/l			15.82	6.5	9.93	7.26	14.84	0	0

Buttons: Print, Save, Refresh, Close

The minimum, maximum, arithmetic mean, as well as other values of interest will be calculated for the selected database parameters.

Once you are finished, you can Save or Print your report.

2.3.1 Saving Reports

Reports can be printed or saved to multiple file formats. The supported file formats are .XLS, .TXT, and .CSV.

NOTE: The pre-defined reports in AquaChem can be saved as .XLS, .TXT, and .CSV. The user-defined reports (e.g. Sample Summary Report created using the Report Designer) can be saved as .HTM or .RTF format.

To save the report,

- ☞ **[Save]** button at the bottom of the report window
- type: a filename for the report
- ☞ Select the file type
- ☞ **[Save]**

2.3.2 Printing Reports

To print any of the reports,

- ☞ **[Print]** button at the bottom of the Report window; or
- ☞ **File** from the main menu and then select **Print**, while the report window is active.

The Report will then be sent to your default Windows printer, or you may select another printer.

- ☞ **[Close]** located at the bottom of the Reports window, to close the window and return to the main AquaChem window.

This concludes the Getting Started chapter.

3

AquaChem Menu Commands

This chapter describes each of the items on the AquaChem menu bar. Similar to the AquaChem Toolbar, the menu is context sensitive and main menu items are available only when the active list window is the currently selected window. In addition, menu items will become grey and inactive if there are no available for the current mode.

In this chapter, you will find information on:

- File Menu.
- Edit Menu
- View Menu
- Filter Menu
- Stations/Samples Menu
- Plots Menu
- Reports Menu
- Tools Menu
- Window Menu
- Help Menu

Following the menu items, you will find a section on the AquaChem database, which covers the database options that are available for your AquaChem project.

The following sections explain the menu commands in detail.

3.1 File Menu

3.1.1 New

The **New** option is only available when no other database is open. In order to activate this option, you must first close any database files which may be open.

The **New** command creates a new empty database based on a database template. The template is an empty database that contains all necessary data tables, parameters, settings, etc., but does not contain sample or station data. When you create a new

database, the database template is copied to the specified filename and location. Choosing a database as a template simply means that you will find the same Lookup-tables, Sample/Station Parameters, and Settings in the database as there are in the selected template.

The AquaChem directory contains two default database templates:

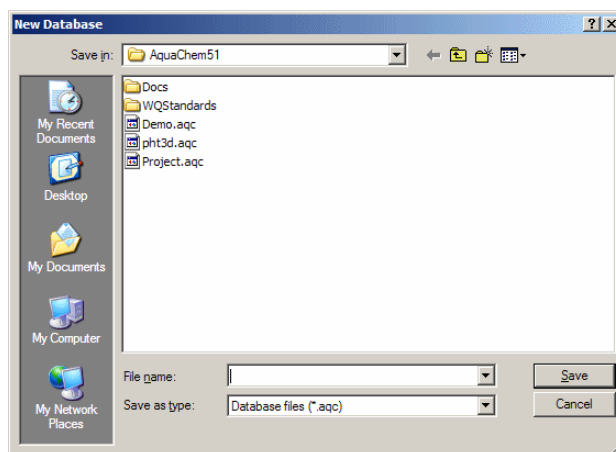
- **Template_Basic.TPL** is normally used to create a new database file.
- **Template_Advanced.TPL** is for more advanced users

AquaChem lets you save and use any valid AquaChem 2011.1 database as a template.

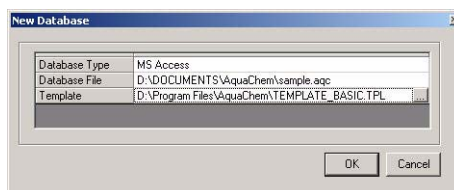
Creating a New Database


Follow the steps below to create a new AquaChem database file:

- Start AquaChem and press **[Cancel]** when prompted to open a database. (Or if you already have AquaChem opened, close the current open database by selecting **File > Close**.)
- Select **New** from the **File** menu, and the **New Database** dialogue will appear as shown below.



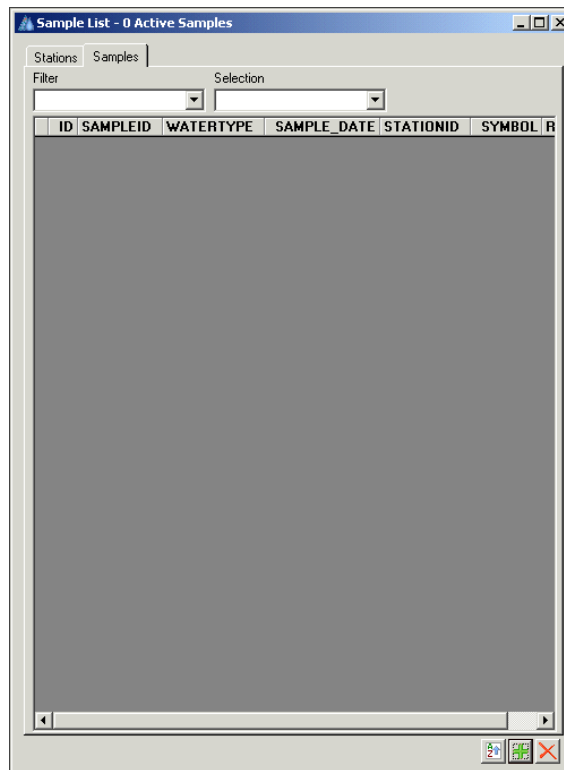
- Browse to the appropriate folder. The default folder is the installation folder *Program Files\AquaChem*.
- Type in the name of the new database, and click **[Save]**.
- You will then see the following **New Database** dialogue.



- Below the database filename, you must specify a **Database Template** to use for the creation of your database. By default, AquaChem will use the **Template_Basic.TPL** file which is included with your installation. This template contains the same data structure as the demo database. When you first start using AquaChem, this will be the only database template file available, and in most cases this template file should be adequate. If you have other database templates available, click the  button to browse and select a .TPL file to be used for your project.
- Once you are finished entering the .database (.AQC) and template (.TPL) name, press **[OK]**.

NOTE: Database filenames should be a combination of letters and numbers only. The filename and directory path should NOT contain spaces, accents, or other special characters.

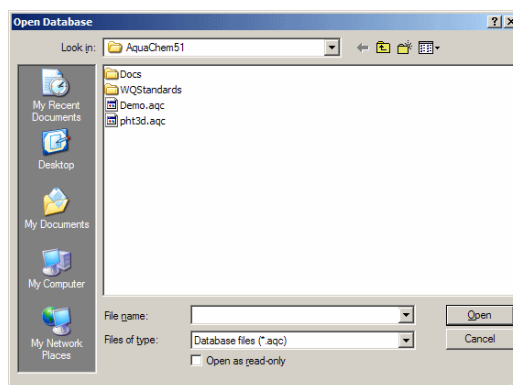
AquaChem will then automatically create a blank database file (with .AQC extension) using the associated database template file containing the data structure for each record in the database. An example is shown below:



If you use the Template_Basic.TPL file, then your data structure may contain some parameters which do not apply to your data set, while others, which you might need, may be missing. Use the **File > Preference > Database Parameters** options to modify the available parameters in the template. These options are explained later in this chapter.

3.1.2 Open

The **Open** command opens an existing AquaChem database. When you select this command, the following dialogue should appear:



You may open an .AQC file from AquaChem v.4.0 or v.5.0. AquaChem no longer supports direct opening of projects from version 3.7 or earlier. To retrieve data from such a project, it has to be exported to a text, MS Access, or MS Excel file and then imported into a new AquaChem database.

NOTE: You may open a database which resides on a network computer. However, only one user should access a database at a time. If you have created your database with an AquaChem version prior to 5.1.20, minor modifications will be applied to your file during opening. You will be informed a backup copy of the original database is created.

After you have selected the desired database, click **[Open]** to load the file into AquaChem. You may also open a database by picking the file from the recent file list, under the **File** menu.

The **Open** database option is only available when no other database is open. In order to activate the **Open** command, you may need to first select **File > Close** to close the opened database.

3.1.3 Close

The **File > Close** command will close the current database. This will allow you to open other databases, or create a new database.

3.1.4 Save Database

Saves the database, or allows you to save the database under a different name and directory. The database file is saved with the extension .AQC.

3.1.5 Save as Template

This option allows you to save the current database settings to a database template (.TPL) file. This new template can be used when creating future databases having the same structure and settings as the currently loaded database.

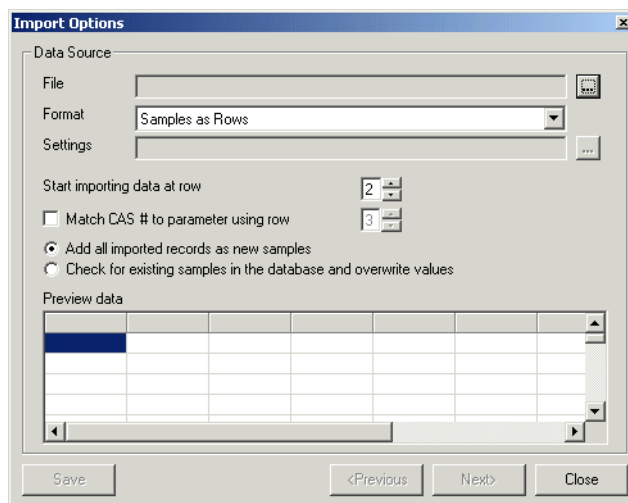
The **Template_Basic.TPL** file is used only in the initial step to create a database. Similar to a word template which you use to create a new word document, the document is completely independent of the template after the document is created. After the database is created, the Template file is no longer needed. You may add, remove, or change parameters and properties; all properties are stored in the .AQC database itself. If you want that new databases use identical settings as your current database, Do not delete or rename the **Template_Basic.TPL** file, since it is the default database template.


The **Save as Template** command is only available when a database is open, and the Active List window is active.

3.1.6 Import

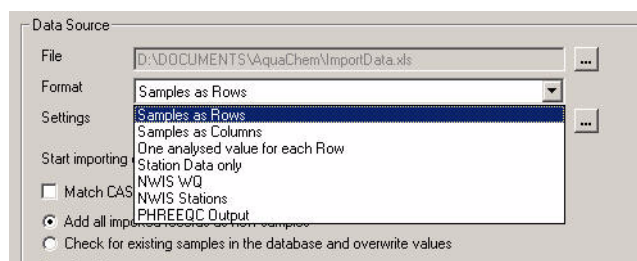
Sample/Stations

The import option allows you to import **Sample** and **Station** data into your AquaChem database. This option is only available when the sample/stations list window is the active window. The Import routine is separated into four simple steps. When you select **File > Import** from the main menu, the following dialogue will appear:



The source data file must be a text file with the data separated by a tab, semicolon, or comma (extension .TXT, .PRN, or .CSV), an MSExcel file (extension .XLS), or an MSAccess 2000 database table or view. When using a character other than a tab (e.g. a comma or semicolon), ensure that this character does not exist in any of the text fields in the file. To load the data file, simply press the  button beside the **File** field.


The file format is specified beside the **Format** field. Use the combo box (as shown below) to select the appropriate file format. The source data file may have the following four formats:



- **Samples as Rows** - each new row represents a new sample, parameters are listed in columns.
- **Samples as Columns** - each new column is a new sample, parameters are listed in rows. The only constraint of this format is that one row has to contain the StationID information for the station for which the sample was collected.
- **One analyzed value for each Row** - each new row contains new measured values. The minimum information that has to be provided in each row is as follows: SampleID (being unique for each Sample), StationID, Parameter name, value, unit (where valid units are: mg/l, ug/l, ng/l). Typically, this format will thus contain further columns with analysis metadata such as MDL (method detection limit), method used for the measurement (Protocol), Precision of measurement, etc. These meta parameters are, however, not required in the import file.
- **Station Data only** - import just the Station Description fields, such as X, Y, Z, Location, Station Name, Well depth, etc. This can be useful if your source data for stations and measurements is kept in separate files. It is then recommended that you import the station data before analysis results. If you select this option, each station must be in a separate row.
- **NWIS Stations** - import station data from the National Water Information System (NWIS) file format (*.txt).
- **NWIS WQ** - import water quality data from the National Water Information System (NWIS) file format (*.txt). **Note:** Be sure to import the corresponding stations before importing WQ data.
- **PHREEQC Output** - import PHREEQC output data as AquaChem samples. For more information on importing PHREEQC output data, please refer to “Importing PHREEQC Output” on page 61.

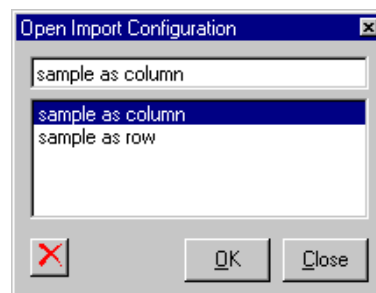
Note: NWIS (<http://nwis.waterdata.usgs.gov/nwis/>) is a large network covering the United States of American, and provides access to water-resources data collected at approximately 1.5 million sites in all 50 States, the District of Columbia, and Puerto Rico.

NOTE: Depending on the format, the first row (or column) must contain headers. Optionally, the subsequent rows may contain the unit or the CAS registry number allowing a more accurate matching of the database and import parameters.

The next line in the **Import Options** dialogue is **Settings**. This option allows you to use previously defined import settings. This is useful for importing datasets with similar formats. If you do not want to use import settings, then leave the settings field blank. If you want to use previous import settings, press the  button to load a list of import configurations:

Select a configuration from the list, then click **[OK]**. If you do not have saved import settings available, then click **[Close]**.

The next step in the **Import Options** dialogue is to specify the row at which to start importing. This option allows you to exclude the first row(s) if it contains header and/or unit data.



AquaChem also provides you the option to identify and match **CAS Registry numbers** (Chemical Abstract Services). If this option is enabled, AquaChem will use the CAS Registry number rather than the parameter name to match parameters from the import file to the database. In order to use this feature, specify the row (or column) of the import file that contains the CAS registry number using the up/down arrows. Also, the registry number must be specified in the AquaChem database parameters. This feature is useful if the import file contains many organic contaminants, which often are named and spelled differently, resulting in an unreliable match between the database and import file when using names only. The CAS Registry number is unique for any registered chemical substances and therefore all parameter matches are effectively realized. It must be noted, however, that the CAS numbers exist only for chemical parameters. Parameters such as TDS, pH, BOD, etc., do not have CAS numbers. If you do not have CAS numbers in your file, then leave this box unchecked.

The import routine has an option to **Check for existing Samples**; this will allow you to add data to already existing samples in your database. When this option is enabled, AquaChem will check for existing SampleIDs in your project database, and if an identical SampleID is found in the database, the import data will be written to the corresponding sample. If the import field includes data which is already in the database, the data in the database is automatically overwritten. This feature is useful, if data for the same dataset has been analyzed by different labs and must be imported from


different source files. If the **Add all imported records as new samples** option is enabled, every record in the import file will create a new sample in the database, regardless if an identical SampleID exists or not. Please note that this may lead to duplicate samples being created in your project database.

Example

To import a data file into your AquaChem database, please follow these steps:

Hint: Before proceeding with the import option, you should ensure that your source file containing your water analysis data is properly formatted. For your convenience, an Excel template is provided with AquaChem, which includes some of the most common sample and station parameters. This file is called **Import_Template.XLS** and can be found in your AquaChem installation folder. If you wish, you may use this file for your data set. Simply enter your data, save the Excel file under a new name and proceed with the import options. Or, you may use the ImportData.xls file as a guide; this file is also included in your AquaChem installation folder.

Step 1: Data Source File

- Select **File > Import** from the main menu.
- Press the  button beside the **File** field to locate the source file.
- Browse to the appropriate folder to locate your source file then click **[Open]**.
Note: If your source file is an Access Database (*.mdb) file, you will be prompted to select the appropriate table or query that contains the source data.
- Your filename should now appear in the **File** field, and a preview of the file should appear in the lower section of the dialogue.
- Select the import **Format** for the source file (Samples as Rows or Samples as Columns, etc.). For this demonstration, the **Samples as Rows** mode will be used.
- Select the row from which to start importing.
- Enable or Disable the option to **Match CAS #'s**.
- Choose the import option to replace or append the existing samples.

Import Options

Data Source

File: C:\Program Files\AquaChem51\ImportData.xls

Format: Samples as Rows

Settings: ...

Start importing data at row: 2

☐ Match CAS # to parameter using row: 3

☒ Add all imported records as new samples

☐ Check for existing samples in the database and overwrite values

Preview data

SampleID	StationID	Geology	Sample_Dat	X	Y	Elevation
MW-1-92	MW-1	sandy-till	8/15/92	533750.2	4814714.5	
MW-1-93	MW-1	sandy-till	6/1/93	533750.2	4814714.5	
MW-1-94	MW-1	sandy-till	6/15/94	533750.2	4814714.5	
MW-1-95	MW-1	sandy-till	7/20/95	533750.2	4814714.5	

Save <Previous Next> Close

- Once you have entered the necessary details in the first import options dialogue, press the **[Next>]** button to proceed.

Step 2: Match Parameters

The next step in the Import data options is to Match Parameters.

The **Match Parameters** frame contains a table listing the Column numbers, the Source text file headers, the Units in the text file, and corresponding AquaChem parameter Internal Key values, as shown in the following **Import Options** dialogue.

Import Options

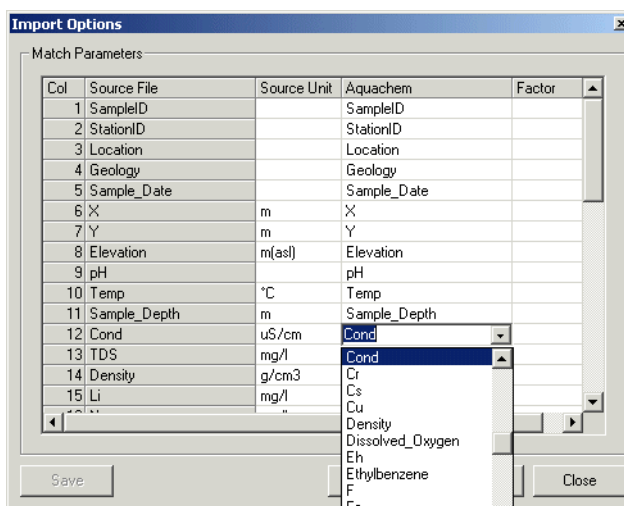
Match Parameters

Col	Source File	Source Unit	Aquachem	Factor
1	StationID		StationID	
2	SampleID		SampleID	
3	X	m	X	
4	Y	m	Y	
5	Elevation	m(asl)	Elevation	
6	Elevation-ScreenTop			
7	Elevation-ScreenBottom			
8	Sample_Depth	m	Sample_Depth	
9	TVOC			
10	Head			
11	pH		pH	
12	Eh	mV	Eh	
13	Temp	°C	Temp	
14	Cond	uS/cm	Cond	
15	TDS	mg/l	TDS	

Save <Previous Next> Close

There is also an option to apply a multiplication **Factor** to the incoming data set. A multiplication factor is needed if the species for the same parameter do not correspond. For example, if Silica is expressed in the import file as mg/L Si and as mg/L SiO₂ in the database. In order to calculate the transformation factor for the respective species you may use the **Species Converter**, available in the **Tools** menu.

In this dialogue, you are required to match up the data in the source file to the appropriate AquaChem parameter labels. If there are blank fields in the AquaChem column, this means that no match parameter was found in the database. This may be due to the lack of this parameter in the database, or that the parameter is spelled differently in the Source file. If the cell stays empty, the respective parameter will not be imported. For example, if your text file had the parameter name **Conductivity**, you must match this up to the AquaChem parameter Internal Key name Cond. Blank fields indicate that AquaChem has not found a matching parameter in the database. If a matching parameter should exist but might be spelled differently than in the data source, you may select it using the steps below:



- Double-click in the grid cell under the **Aquachem** column and a combo box will appear listing all of the available parameters in your AquaChem database.
- Select the appropriate parameter to match to the import **Source File** column. This will correctly link the parameters in the source file to the parameter names in the AquaChem database template.
- If the parameter to be imported is not included in the database, it may be created prior to importing. Select **<create>** from the available list. An AquaChem database parameter will be created for every Import parameter that has been mapped to **<create>** when you click the **[Next]** button. You will be prompted to enter the required information for each parameter to be created in such way.

- Repeat this step for each parameter in your source file which has not been correctly mapped to an AquaChem database parameter.

Hint: If the field names in the file are identical to the field names in the AquaChem database template file, the parameters will be mapped automatically to the AquaChem column. Therefore, if you format your source file correctly prior to importing, you can save some time in this step.

AquaChem allows you to import concentrations in the units which may be different from the ones specified for this parameter in the database. These values will be converted to the units used in the database during the import. Unless otherwise specified, the units used for that parameter will be identical to those specified in the database parameters options (under **File > Database**). To change the default units, double-click in the corresponding grid cell under the **Units** column and you will see a selection of available units (g/L, meq/L, mg/L, mmol/L, mol, ppm, umol/L, ug/L). Select the desired unit, and repeat this for other parameters as needed. The values will be converted into the project units.

Once you have mapped all units and parameters, press the **[Next>]** button to proceed.

If you have at least one Import parameter mapped to <create>, the following dialogue will load:

	Value
Internal Key	SiO2
Form Label	SiO2
Description	silica
Formula	SiO2
Formula Weight (g/Mol)	60.0843
Solubility	0
Valence	0

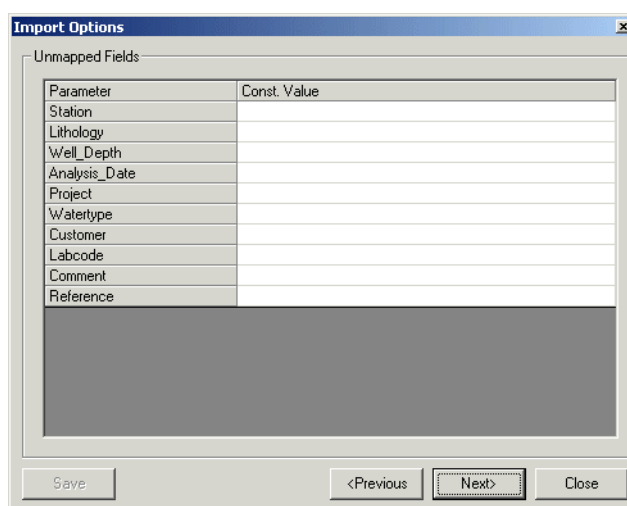
Fill in the appropriate data and click **[OK]**. Make sure you select the appropriate parameter type using the **Type** combo box at the top of the dialogue. You may later edit these options later on in **File>Preferences>Database Parameters**. Click **[Select]** to copy predefined information (CAS number, formula) from a list of predefined parameters. The most important value that needs to be provided is the parameter name. Other values can be specified after the import. If more than one parameter was mapped to <create>, AquaChem will present a new dialogue for each parameter, which was marked with the <Create> flag.

Once the last parameter was created, the dialogue will close and you will move to the next step of the Import routine.

If the data does not appear to be correct, press the [**<Previous**] button to return to the previous step.

Step 3: Unmapped Fields

The next dialogue contains a list of the AquaChem parameters which were not matched up to parameters in the source file (ie. **Unmapped Fields**).



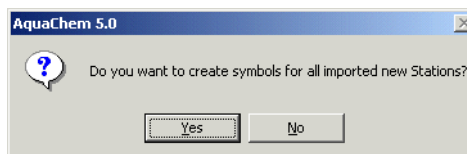
The 'Import Options' dialog box, 'Unmapped Fields' tab, is shown. It contains a table with two columns: 'Parameter' and 'Const. Value'. The parameters listed are Station, Lithology, Well_Depth, Analysis_Date, Project, Watertype, Customer, Labcode, Comment, and Reference. Below the table is a large empty text area. At the bottom are buttons for 'Save', '<Previous', 'Next>', and 'Close'.

Parameter	Const. Value
Station	
Lithology	
Well_Depth	
Analysis_Date	
Project	
Watertype	
Customer	
Labcode	
Comment	
Reference	

This dialogue allows you to enter constant values. For example, if all of the imported samples belong to the same project or to the same location, but this information is not included in the data, you may fill the respective fields automatically with the specified value during this step of the import. For example, you may read the data from a file called MW-1.xls indicating that all samples were taken from well MW-1. In this case, you would enter "MW-1" as a constant value for location. To skip this step, leave these fields blank.

To proceed to the last step in the import routine, press the [**Next>**] button. To return to the previous step, press the [**<Previous**] button.

Upon proceeding, you will then see the following prompt:



The 'AquaChem 5.0' dialog box is shown. It contains a question mark icon and the text 'Do you want to create symbols for all imported new Stations?'. Below the text are 'Yes' and 'No' buttons.

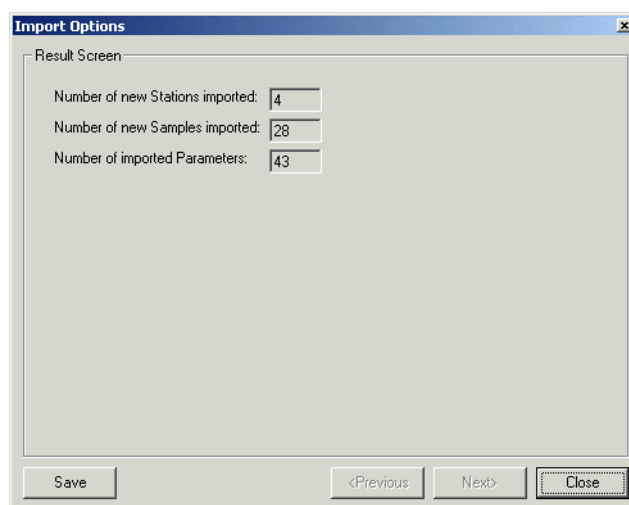
This option allows you to automatically create one symbol for each unique station in the Station symbol group and assigns this symbol to the appropriate samples.

☞ **[Yes]** to accept this feature, or **[No]** to ignore this option.

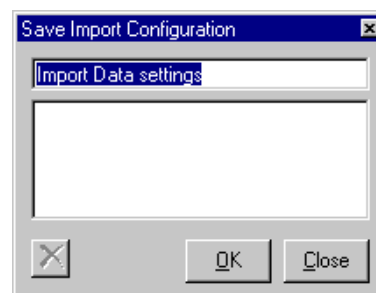
Step 4: Result Screen

The last dialogue in the Import options is a summary of the results of the data import.

The **Result Screen** dialogue shows the Number of new Samples and Stations that were successfully imported, and the Number of parameters that were successfully mapped and imported.



Before closing the import routine you can save your import settings by clicking the **[Save]** button. This option allows you to save the import settings (file format, mapped parameters, units, factors, and constant values) for future use. When you click the **[Save]** button you will be prompted with a **Save Import Configuration** dialogue as shown on the right hand side. Simply enter a name for the import configuration, click **[OK]** and then **[Close]**. After successfully importing the data, you will return to the main database window showing your active samples and stations. This concludes the necessary steps for importing data. Once you are finished, press the **[Close]** button.



Importing PHREEQC Output

AquaChem can import files are that in PHREEQC selected output format. This is useful if you generated a number of simulations for example by mixing two samples in different proportions and you wish to save the simulations as AquaChem samples so that you can use the program's plotting and reporting capabilities to analyze the results. Prior to this import, the database must include at least one stations, and all parameters in the PHREEQC output file must exist in the database parameters within AquaChem. To confirm that the required database fields exist in the database, or to add fields to the database, navigate to the **Database** settings by selecting **File > Preferences** from the main menu, and then selecting the **Database Parameters** option.

To import the PHREEQC output file, follow the steps below:

Selecting the Data Source

From the main menu, select **File > Import > Sample Stations**

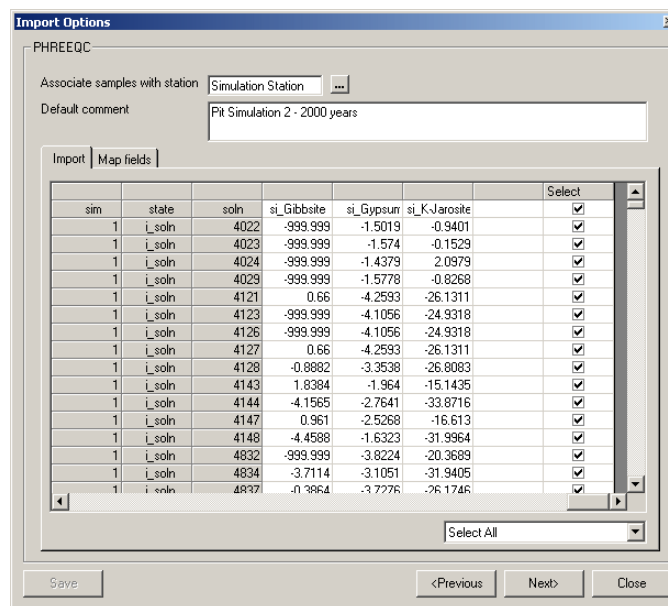
The **Import Options** dialog box will appear on your screen.

sim	state	soln	dist_x	time	step	pH	pe	reaction
1	i_soln	4022	-99	-99	-99	7.75	4.25	.
1	i_soln	4023	-99	-99	-99	7.04	4.96	.
1	i_soln	4024	-99	-99	-99	7.57	4.43	.
1	i_soln	4029	-99	-99	-99	8.24	3.76	.
1	i_soln	4121	-99	-99	-99	6.855	0.0712245	.
1	i_soln	4123	-99	-99	-99	6.75	0.0712245	.
1	i_soln	4126	-99	-99	-99	6.75	0.0712245	.
1	i_soln	4127	-99	-99	-99	6.855	0.0712245	.
1	i_soln	4128	-99	-99	-99	5.76	0.0712245	.
1	i_soln	4143	-99	-99	-99	6.255	0.0712245	.
1	i_soln	4144	-99	-99	-99	3.25	0.0712245	.
1	i_soln	4147	-99	-99	-99	7.63	0.0712245	.

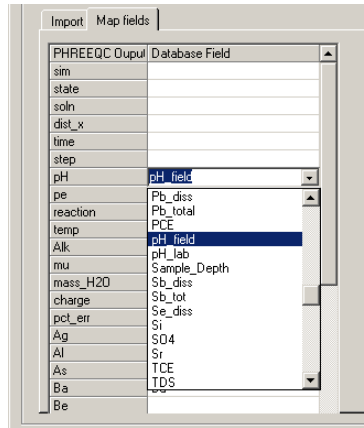
- Click the [...] button located beside the File field, and select the desired PHREEQC output source file.
- From the **Format** combo box, select the **PHREEQC Output** option.
- Select which row to start importing data from
- Choose to add the imported records as new samples, or overwrite existing samples in the database using the appropriate radio button.
- Click the **[Next]** button.

Assign Station & Map Fields

The following dialog will appear on your screen:



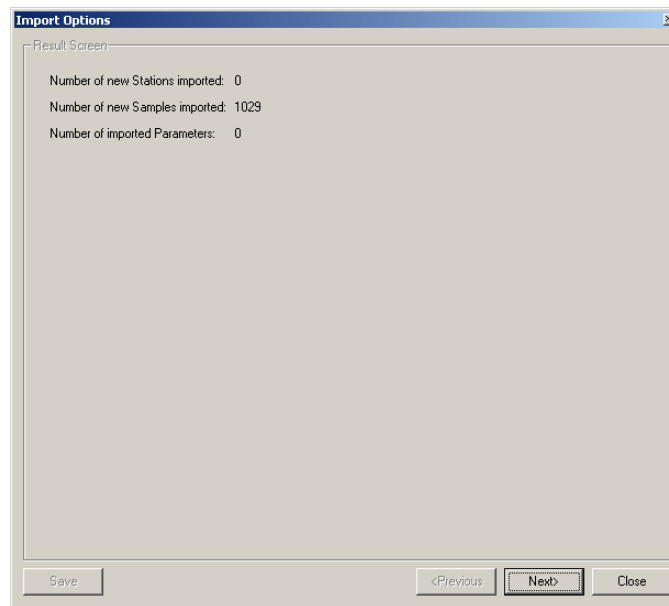
- Click the [...] button, and select the associated station from the list of stations.
- Type a comment describing the simulation results (optional)
- Under the Import tab, using the horizontal scroll bar, scroll across to the far right of the preview table. In the far right column, labeled “Select”, select the desired samples to include in the import. You can use the combo box located below the table to make automatic selections from the table, e.g., Select All, Select Solutions only, etc.
- Click the Map Fields tab. Map the PHREEQC output fields to the appropriate fields in the AquaChem database. **Note:** AquaChem will automatically map fields that have the same parameter name.



- Click the [Next>] button

Viewing Import Results

The data will then be imported. Once finished, the results screen will appear displaying the number of samples imported (shown below).



- Click the **Close** button to close the **Import** dialog.

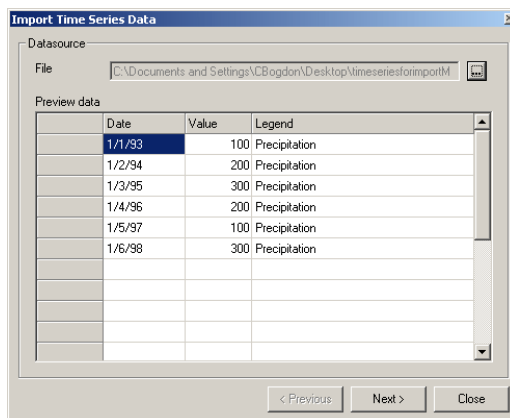
Time Series Data

AquaChem allows you to import non-chemical time series data, *i.e.*, water levels, precipitation, evaporation, etc., from Text Files (.PRN, .TXT, .CSV, .OUT), Microsoft Excel spreadsheets (.XLS) and Microsoft Access databases (.MDB).

Once imported, the time series data can be plotted independently and compared with chemical data on time series plots.

To import non-chemical time series data, select **File > Import > Time Series Data**, from the AquaChem main menu.

The following dialog will appear on your screen:



Selecting the Source File

Select the [...] button beside the **File** field to specify the desired source file. A preview of the data will show in the Preview Data table below. Please consider the following when importing time series data:

- The source field must contain the following columns: **Date**, **Value** and **Legend**. The **Date** column contains the dates for each measurement in MM/DD/YYYY format. The **Value** column contains the measurements of an attribute, e.g., precipitation, water levels etc. The **Legend** column contains a descriptive label for each row which will be used for showing on plot legends, e.g., Precipitation, Water Levels, and for distinguishing groups of data within the same time series table.
- Each column in the source data should have a header row, e.g., Date, Value, Legend.
- Any rows that contains a blank value for one of the above columns will not be imported.

- When importing data from an Access database (*.mdb), you will be prompted to select the desired Access table or query.

Click the **[Next]** button to proceed to the next step in the import process.

Matching Columns

Specify a **Series Name** by selecting the green plus and a brief description of the time series data. Please note that the **Series Name** will be used to identify the time series when plotting the data on a time series plot. It is recommend that you create a new table for any distinct data type, such as water levels, participation, pumping etc.

The next step involves mapping the fields in the source data to the required fields in the AquaChem database, and assigning the appropriate data type to each field.

Source Field	Match
SampleID	
Sample_Date	
Analysis_Date	
Project	
Waternumber	
shortWaternumber	
Comment	
Reference	
Quality	

Note: To import Time Series data, a Series Name must be specified and each Mapping type (Date, Value and Legend), must be mapped to a row.

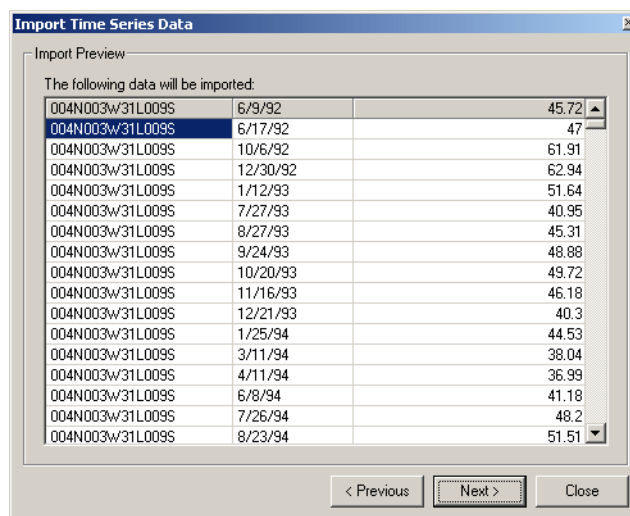
Under the **Match** column, select the appropriate destination field for each field in the source data. You must specify a field for Value, Date and Legend. For example, the date field in the source data should be matched with the **Date** field, and the measurement field should be matched with **Value** field. The **Legend** field should be assigned to a column in the source data that contains a descriptive label, e.g., Water Level. This label will be displayed on the time series plot legend.

Note: Each time series may only have one **Value** field (measured attribute). Additional attributes must be imported as a separate time series.

Once the fields have been mapped, select the **Next>** button.

A warning message may appear if any rows in the source data are deemed invalid (see below).

A preview of the source data will appear (shown below).



The following data will be imported:		
004N003w31L009S	6/9/92	45.72
004N003w31L009S	6/17/92	47
004N003w31L009S	10/6/92	61.91
004N003w31L009S	12/30/92	62.94
004N003w31L009S	1/12/93	51.64
004N003w31L009S	7/27/93	40.95
004N003w31L009S	8/27/93	45.31
004N003w31L009S	9/24/93	48.88
004N003w31L009S	10/20/93	49.72
004N003w31L009S	11/16/93	46.18
004N003w31L009S	12/21/93	40.3
004N003w31L009S	1/25/94	44.53
004N003w31L009S	3/11/94	38.04
004N003w31L009S	4/11/94	36.99
004N003w31L009S	6/8/94	41.18
004N003w31L009S	7/26/94	48.2
004N003w31L009S	8/23/94	51.51

Rows that cannot be imported will be highlighted yellow. A row will be deemed invalid if it meets one of the following criterion:

- The row contains empty cells
- The data type of the source field and target field do not match. For example, if you assign the Date field as a Text data type in the mapping step instead of a Date data type.

Click the **Next>** button to continue.

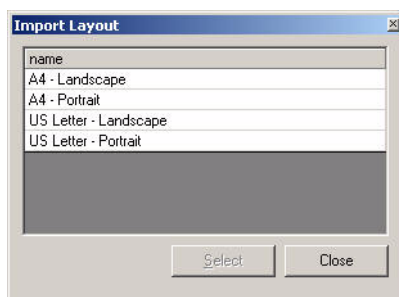
The **Results** dialog will appear, displaying the number of imported records. Click the Close button to complete the import process.

The imported time series can now be viewed in **AquaChem** by selecting **Tools > Lookup Tables > Time Series** from the main menu.

For information on showing time series data on time series plots, please refer to “Time Series (Multiple Parameters)” on page 241.

Page Template

This import option allows you to import Report Page Templates from other AquaChem projects. First specify the AquaChem project file (*.aqc) you want to import a Page Template from. Then select from the list of Page Templates which layouts you wish to import (you can select more than one) and choose the Select button.



You will find the imported Page Templates in the Template Designer.

Plot Layout

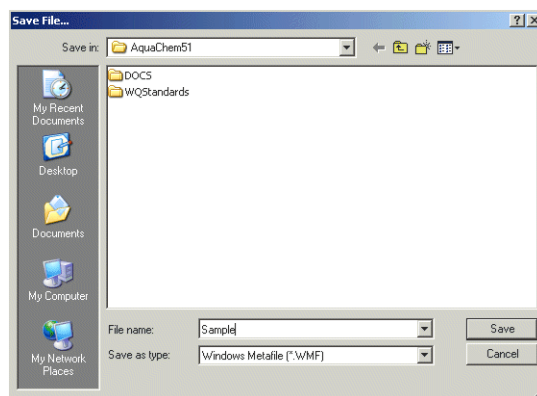
This import option allows you to import Plot configurations from other AquaChem projects. First specify the AquaChem project file (*.aqc) you want to import a Plot Layout from. Then select from the list of Plot Layouts which you wish to import (you can select more than one) and choose the Select button.

3.1.7 Export

The **Export** option allows you to export the data or plots for use in other applications such as **Visual MODFLOW** and **Hydro GeoAnalyst**. The **Export** command is only available when the Active List window or a graph window is currently active.

Image

When any graph window is active, the **File>Export>Image** option is available. Clicking this option will load the following dialogue:

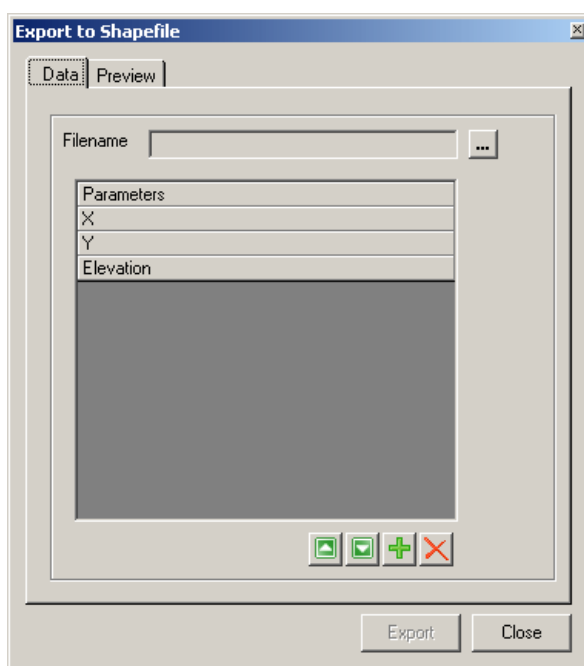



You have the option of saving the image in .WMF (Windows metafile) format.


ESRI Shapefile

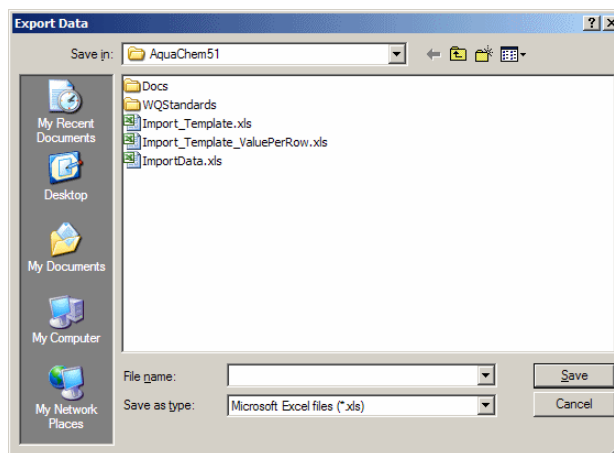
This option is available only when you have the Active Samples list or a Map plot active and it allows you to export the data as an ESRI shape file (.SHP).

If the Active Samples list active, selecting this option will load the following dialogue:



Use the  button to add parameters to export (X and Y coordinates are selected by default and cannot be removed).

To create the destination file, click the  button beside **Filename** field. The following dialogue will load:



Type in the name of the destination file that you wish to create and click **[Save]**. You will be returned to the previous dialogue and the File name field will now be filled with the appropriate address.

Click **[Export]** to export this point theme shapefile. The values for all specified parameters will be included for every point in the attribute table.

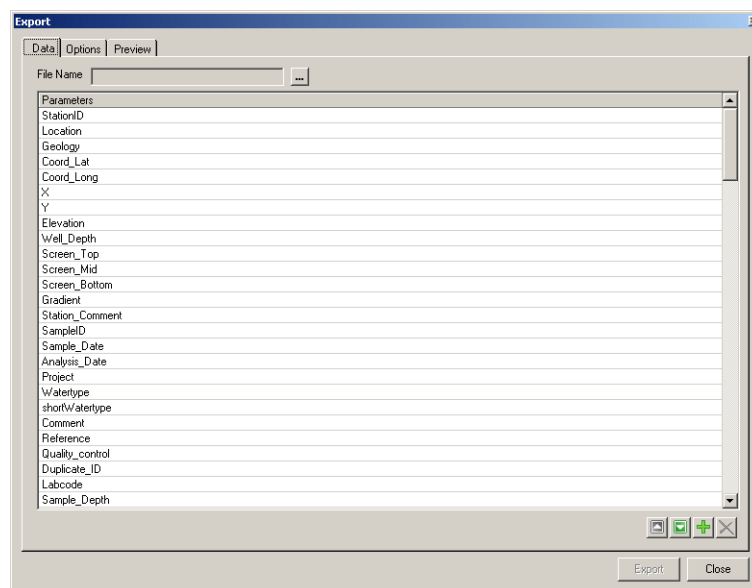
If a Map plot is active, selecting the **File>Export>ESRI Shapefile** option will produce a point or a polygon theme shapefile depending on the currently active symbol type. If the map shows a plain symbol, scaled symbol or a pie plot, then a point theme will be created holding all the chemistry per point in the attribute table. For example, for a pie plot - all parameters; for a plain symbol - just StationID. If the map plot shows a Stiff or Radial diagram per sampling point, then the respective diagrams are exported as polygons, along with the current symbol label, and the data attributes. Since these polygons are defined in map coordinates, you should optimize the symbol size in AquaChem according to the final map scale in Hydro GeoAnalyst or ArcView.

Type in the name of the destination file and click **[Save]** to export the file.




Data

This menu item is available when the Active Samples list or a plot is active.

To export your sample/station data, select **File>Export>Data** while the Active Samples list is active. The following dialogue will load:



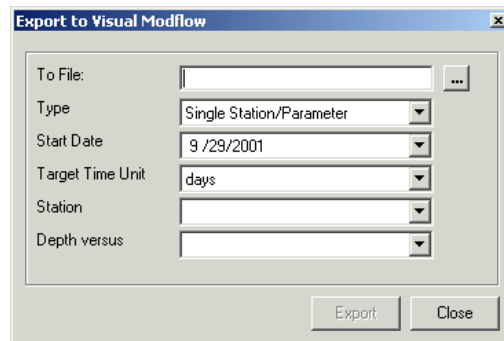
Data may be exported in 2 formats: **Samples as rows** and **One parameter per row**. These options are available under the **Options** tab. The **One parameter per row** format means that for each parameter the stationid, sampleid, value, unit, analysis comments, precision, MDL will be exported. This allows you to export the complete information into an export file. However this format is less convenient if data is to be used for plotting, statistics, etc. In the case of the standard **One parameter per row** format, only the concentration value for each measurement result will be written to the export file.

- By default, all the parameters are selected and will be exported.
- To remove parameters from the export, highlight the parameter you wish to remove and click the  button.
- To add parameters to the list, click the  button and select from the parameters list that loads.
- You may choose the parameter names or the parameter labels to be used as headers in the destination file. Parameter labels are a good choice if you wish to create a table or a presentation. Parameter names should be considered if the exported table should be re imported back into a different AquaChem database, as the parameter names will match automatically.
- Click the  button beside the File name field to create the destination file. The destination file can be of type .PRN, .TXT, .CSV, and .XLS.
- You may view a Preview of the exported data under the **Preview** tab.
- Once you are finished, click the **[Export]** button to generate the file.

You may also export data that was used to generate a specific plot. This feature is useful if you wish to use this data in another plotting program. To export plot data, select the plot you wish to export the data for and then select **File>Export>Data**. Once a window containing the pertinent data loads, you have a choice of printing the data or saving it as a .TXT, .CSV, or a .XLS file.

MODFLOW

The **Export > MODFLOW** option will export your current sample and station data to a space-delimited text file. The data can then be used for the input (concentration observation well data) of a Visual MODFLOW groundwater model.



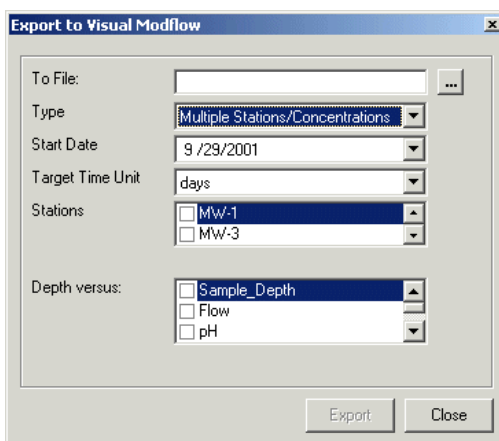
To export the data:

- Select **File>Export>Visual MODFLOW** from the main menu.
- Click the **...** button beside **To File** to generate the destination file. This file can be a .TXT, .CSV, or .XLS file.
- Select the **Type** of data to export. The default is Single Station/Parameter, however you may choose Multiple Stations/Concentrations as well. See below for a more extensive description of the second option.
- Select the **Start Date**
- Select the **Target Time Unit**
- Select the **Station** to export
- Select the parameter to export
- Click **[Export]** to export the file

Each line in the generated file will have a time measurement and an observed concentration for a single parameter. The start date is used to calculate the time for the first observation.

NOTE: The **[Export]** button will become active only after the necessary options have been specified.

- If you choose **Multiple Stations / Concentrations** format, then the dialogue shown below will appear:



- Select the **Start Date**
- Select the **Target Time Unit**
- Select the stations you wish to export by checking the boxes beside them
- Select the parameters you wish to export by checking the boxes beside them

This format is only practical if a **Depth** parameter value is available in your database. After selecting all desired information, the observation name is concatenated with the station code and depth in the resulting destination file.

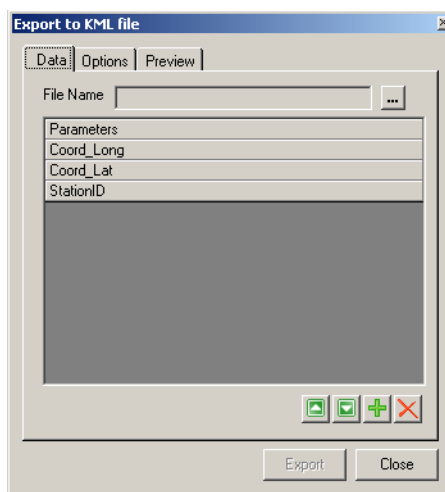
Once you have specified the necessary settings, click the **[Export]** button to proceed and generate the export file. Press the **[Close]** button to close this dialogue and return to the main AquaChem window.

KML File

This option allows you to export **Station** coordinates and associated parameter data to .KML file, a file format supported by various geobrowsers, such as Google Earth™. When a KML file is imported into Google Earth™, each station is projected on a 3D globe as a single point symbol. Associated sample data may be viewed in a pop-up information box for each station.

Note: .KML format requires station locations in geographic coordinates, e.g., latitude and longitude. Therefore, if your station coordinates are currently expressed in a projected coordinates system (e.g., UTM), you must convert them to geographic coordinates (e.g., Lat/Long). This can be easily done using the AquaChem UTM conversion tool (see “UTM Conversion” on page 342.)


To export data to KML file format, select **File > Export > KML File** from the main menu. The **Export to KML File** dialog will appear on your screen.



The **Export to KML file** dialog consists of three tabs: **Data**, **Options**, **Preview**.

Data

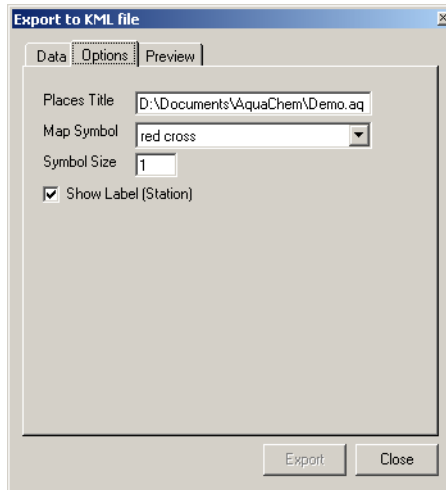
Select the [...] button located beside the **Filename** field. The **Export Data** dialog box will appear, where you can specify the desired destination folder and file name for the .KML file.

Use the  button to add parameters to the .KML file. When selected, the **Parameters** dialog box will load, where you can select one or more parameters in the AquaChem database. Please note, Coord_Long, Coord_Lat and Station ID are required parameters, and are selected by default. These parameters cannot be removed.

If you wish to export sample data, be sure to add the Sample ID parameter.

Options

The **Options** tab provides several export options for displaying data in Google Earth.



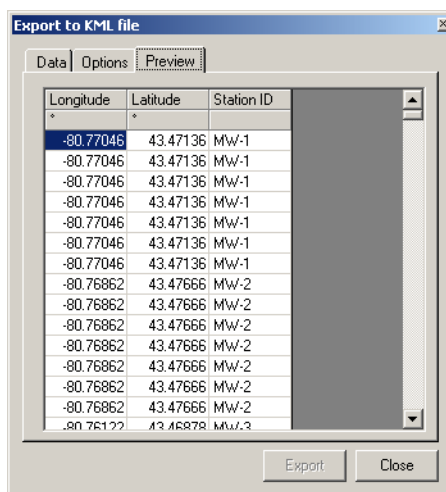
Choose the desired station symbology from the **Map Symbol** combo box. When the KML file is opened in Google Earth, each station will be displayed on the map using the specified symbol.

The **Symbol Size** text box allows you to specify the size of the Map Symbol, when displayed in Google Earth. Enter a value between 1 (smallest) and 10 (largest). Please note, the map symbols can be further customized using various options provided by Google Earth itself.

The **Show Label** allows you to enable or disable station labels. If this checkbox is empty, by default station names will not be displayed as labels in Google Earth. If this checkbox is “checked”, a station name label will appear beside each station.

Preview

The **Preview** tab provides a preview the data that will be exported to the KML file. The data displayed in the Preview tab is read-only.



Once you have selected the desired parameters, and set the appropriate options, click the **Export** button to generate the .KML file. Upon exporting, you will be prompted to open the file in Google Earth™. If you have Google Earth™ installed, click the **Yes** button to open the .KML file. If you do not have Google Earth™, click the **No** button. The file can be imported manually into Google Earth™ by selecting **File > Open** from the Google Earth™ main menu.

PDF File

This option allows you to export Plot reports to PDF file format.

Note: This option is only available when the Print Options window is shown on your screen.

To export to PDF file format:

- Create the desired plots (**Plots > New**), or open a saved plot configuration (**Plots > Open Configuration**).
- Select **File > Print**, from the main menu.
- The **Print Options** window will appear on your screen. Configure the plot report as desired using the available options.
- With the **Print Options** window open, select **File > Export > PDF file** from the main menu.
- The **Export to PDF** file dialog will appear on your screen. Specify the destination folder and the desired file name, and then click the **Save** button.

Please note that you must have a version of Adobe Acrobat reader installed in order to view exported PDF documents.

3.1.8 Remote Database

Link to Remote Database

This option allows you to link remote **Hydro GeoAnalyst (HGA)** databases to the AquaChem interface through an Open Database Connectivity (ODBC) connection.

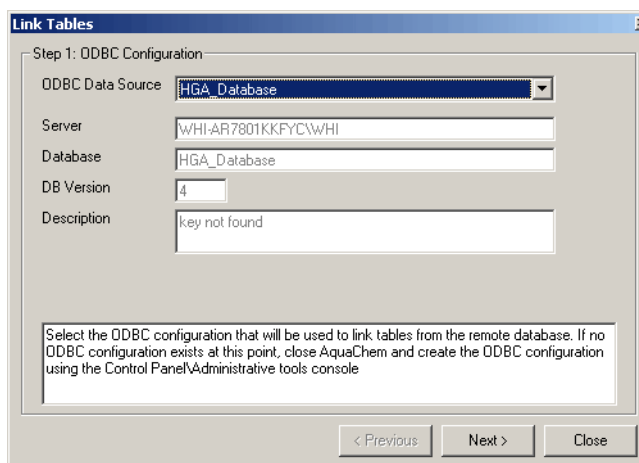
Requirements

The following is required for setting up an HGA-AquaChem linked database:

- An ODBC connection must be configured for the HGA database. For instructions on how to set up an ODBC connection, please see “Appendix B: Configuring an ODBC Connection” on page 415
- Only projects that use the SWS Environmental database template can be linked with AquaChem. If your project uses a different database template, please contact SWS for more information.
- If your database was created using HGA 2009.1 or earlier, you must make the following changes to your database using the HGA template manager.
 - In the Parameter_Result table, make the **sample_id** and **chemical_name** field a Primary Key.
 - In the Parameter_Result table, change the **Data type** of the “**Outlier**” field to “**Double**”.
 - In the Parameter_Result table, rename the “**comment**” field to “**result_comment**”.

Follow the steps below to link your HGA database to the AquaChem interface.

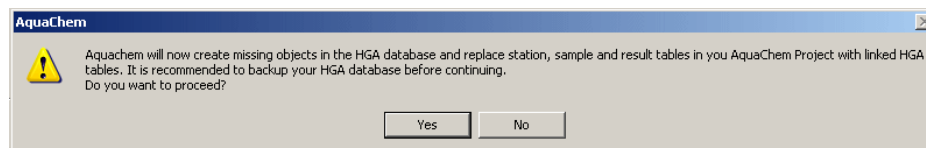
- [1] Launch AquaChem
- [2] Create a new empty AquaChem database. Please see “New” on page 47 for more information on creating a new empty database. It is important that a new AquaChem database is created as once the tables are linked, the AquaChem tables are deleted and data may be lost.
- [3] From the main menu, select **File > Remote Database > Link to Remote Database**. The following dialog will display:



From the **ODBC Data Source** combobox, select the ODBC configuration that will be used to link tables from the remote database. If your HGA database does not appear in the combobox, review the ODBC connection settings to ensure it is set up correctly.

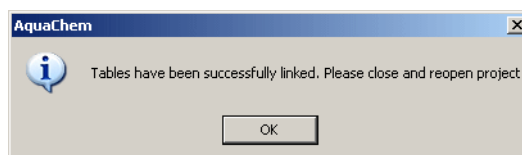
Once the database is selected, its corresponding server information will be shown in the **Server** and **Database** field.

- [4] Click the **[Next]** button to begin the automated linking process. The following warning message will show:



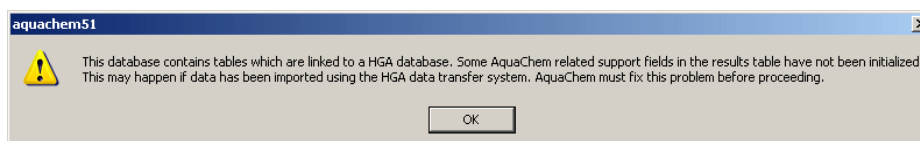
Before proceeding it is highly recommended that you back up your HGA database.

- [5] Click the **[Yes]** button to continue.
 [6] If the linking is successful, the following message will show:



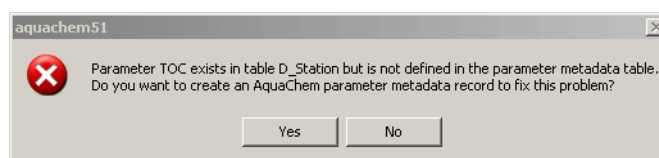
- [7] Click the **[Ok]** button
 [8] Close the **Link Tables** dialog by clicking the **[Close]** button
 [9] From the main menu, select **File > Close** to close the current AquaChem project

- [10] From the main menu, select **File > Open** to reopen the AquaChem project. Locate the project file in the **Open Database** dialog and then click **[Open]**
- [11] Upon reopening the project, you may receive the following message:



Before continuing, AquaChem needs to initialize some fields in the HGA database. Select the **[OK]** button to continue.

- [12] In some cases the following message may appear:



This message will show when there are duplicate field names in the station table and the sample table. If your HGA database already contains chemistry data, it is likely that this message will show. Simply click **[Yes]** and AquaChem will fix the problem.

Now that the tables have been successfully linked, any changes made in one application, will show in the other. For example, if you modify a station name in the AquaChem station table, this change will also show in the HGA station list, and vice versa.

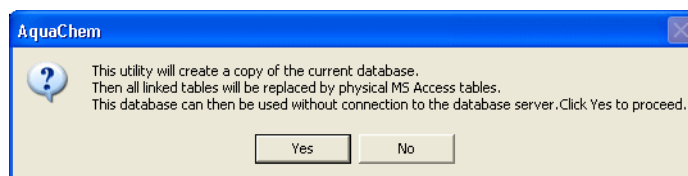
Please note that AquaChem is now accessing data from a SQL Server database. This means that the SQL Server must be running and you must be connected to the server in order to access its data.

Note: After linking, some of the linked fields, e.g., sample_id, may not show as visible in the station and sample tables in AquaChem. To make them visible, go to **View > Options** and add the fields in the sample/station list options.

Import Linked Tables

If you are connected to a HGA linked SQL Server database, you can use this option to convert the linked tables to a MS Access database format (*.aqc). This allows you to use the database without a connection to the SQL Server.

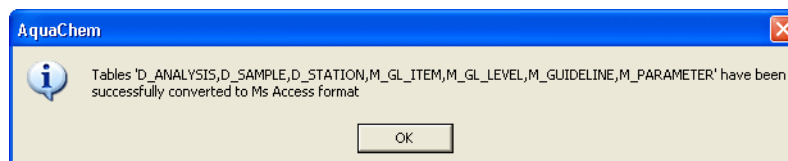
When this option is selected, the following message will be shown:




Click the **[Yes]** button to proceed.

A save dialog will display prompting you to specify a name and location for the database. Once specified, click the **[Save]** button.

When the tables have been converted to MS Access database format, the following message will be shown.



3.1.9 Print

The **Print** command will load the print options for the current AquaChem window or dialogue. The print icon  in the toolbar serves the same purpose. The **Print** option is only available when one of the following windows is active:

- Plots
- Tables (Spreadsheet View), or
- Reports

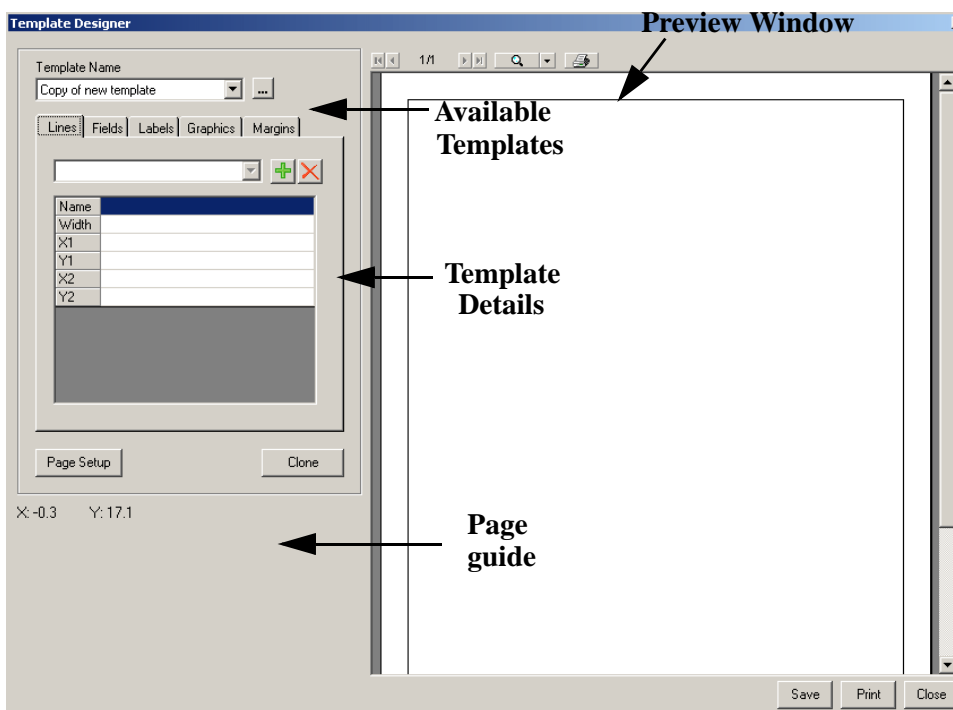
Please refer to Chapter 4: Printing and Exporting Plots for more details.

3.1.10 Template Designer

Use the Template Designer to customize the pre-defined printing templates shipped with the AquaChem demo database, or to create new ones.

Template Designer Interface

When you select **Template Designer** from the **File** menu, the following window will appear:



The **Template Designer** dialogue has three main components:

- [1] **Available Templates** in the upper-left corner containing the **Template Name** and a combo box listing the available templates.
- [2] **Template Details** is located immediately below the **Template Name** field. This includes the controls and tools for designing templates, such as:
 - **Lines,**
 - **Fields,**
 - **Labels**
 - **Graphics,** and
 - **Margins**
- [3] **Preview Window** on the right side of the dialogue, containing a preview of the Template as content is added or modified.

The options in the **Template Designer** allow you to:

- Select the page size, orientation, and margins
- Draw lines and borders
- Define and position text block descriptors
- Position user-entry fields
- Add headers and footers, and

- Position and size bitmap graphics for a company logo
- Adjust the margins of the plot display area

The following section provides more details on the functionality of the template designer.



Template Designer Controls


The main controls and tools for the **Template Designer** are on the left side of the main dialogue (as shown in the figure below).


At the top of this frame, you will see a list of the available templates; you can select a different template from the combo box. AquaChem includes the following pre-defined templates:

- US Letter - Portrait
- US Letter - Landscape
- A4 - Portrait
- A4 - Landscape



In the **Template Designer** dialogue:

- Click  button to add a new item to the template.
- Click  button to delete the currently highlighted item.

Note that the buttons are common to each tab. For example, if you are in the **Lines** tab and you press the  button, a new border line will be added. You can then proceed to enter the appropriate properties for this line in the fields below.

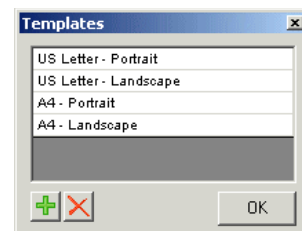
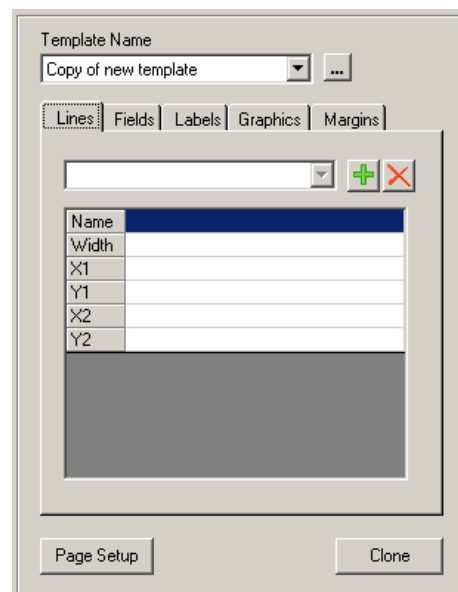
The  button in the **Template Designer** dialogue will load a **Template Name** dialogue as shown on the right. It displays the full list of templates currently available and will allow you to create and name new templates.

In the **Template Name** dialogue:

- Click  button to create a new template
- Click  button to remove an existing template

To rename a template simply place your mouse cursor in the appropriate field, then enter a new text label. Once you are finished in this dialogue press **[OK]**.

When positioning a line, field, or graphics, the X, Y page coordinates displayed in the lower left section of the main **Template Designer** dialogue can be helpful. These

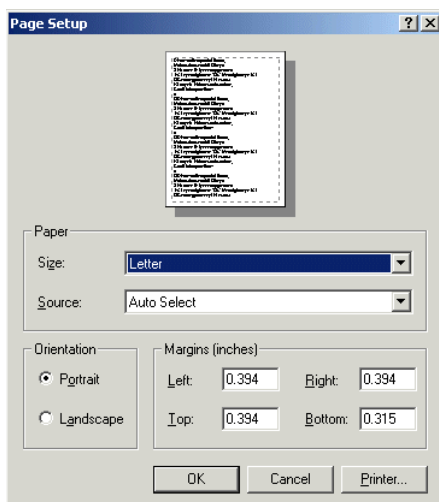


coordinates correspond to your mouse cursor location and can help you in the placement of the start and end points.

The **[Duplicate]** button will duplicate the currently selected template. The duplicate template will be named **Copy of TemplateName**, and the duplicate template will have the same layout and characteristics as the originally selected template. If it is your first time creating or customizing a Print Template, it may be easier to modify a duplicate copy of an existing template, than to start from the beginning. Please note that each Template must have a unique name.

NOTE: Although you have the option of modifying the default template designs, it is strongly advised that you either duplicate the selected template, or create a new blank template. This avoids damaging or losing the original template designs and ensures you will always have a standard set of default Print Templates.

When you press the **[Page Setup]** button, the following **Page Setup** dialogue will appear:





In this dialogue, in the upper **Paper** section you can define paper **Size** and **Source**.

In the lower **Orientation** section, you can define paper orientation and page **Margins** for the template. The page margins will correspond to the location of the page border. The page dimensions inside the margins will be used to automatically locate and size the selected plots.

You can also access your standard Windows Printer options by pressing **Printer** button. Once you are finished, press **[OK]** to return to the previous dialogue.

Creating New Templates - Example

The following section describes the options of the **Template Designer**, and how it can be used to build and customize your own templates. The brief instructions below tell you how to create a basic printing template, and how to customize this to your own preferences.

- Click  button to load the list of Templates.
- Click  button to create a new template.
- Enter a name for the new template (e.g. “Sample-Letter” for these instructions).
- Press **[Close]**.
- Choose this template from the combo box.


You will then see a new blank Print Template in the preview window. By default, all new templates include a border and default page settings.

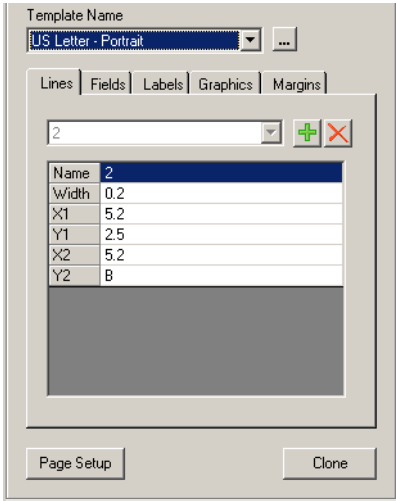
Creating Lines

The first tab in the template designer dialogue is **Lines**. This tab allows you to create the Print Template form by drawing lines indicating the borders of the template Title Blocks. Each line has a **Name**, **Width**, starting coordinates (**X1**, **Y1**), and ending coordinates (**X2**, **Y2**). The coordinates units are in cm. The coordinates are measured from the origin, which is the bottom left corner of the page.

In the **Border** frame you can show/hide the page border using the check-box, as well as set the border width.

To create a new Line:

- Click  button and a new Line will be created with the name **new Line#**.
- Double-click in the **Name** field and enter a unique name for the new line (e.g. Line2).
- Enter a line **Width** 0.2.
- Enter the desired start and end coordinates for the new line (**X1**, **Y1** represents the starting point of the line, and **X2**, **Y2** represents the end point of the line). The bottom-left corner of the Preview Window serves as the origin for the Template designer.



To edit an existing line, select the line from the list and enter the new coordinates. For example, if you want to start a line at a point located 1.5 cm from the bottom of the page and 1.5 cm from the left side of the page, you would enter a line with **X1 = 1.5** and **Y1= 1.5** (cm). Remember that you can always use the mouse pointer to check the page coordinates, as shown on the left side of this dialogue.

If you want to enter line locations to coincide with margin locations, enter the following items for the X or Y coordinate:

Enter “**L**” to have a line coordinate equal to the **left** margin location.

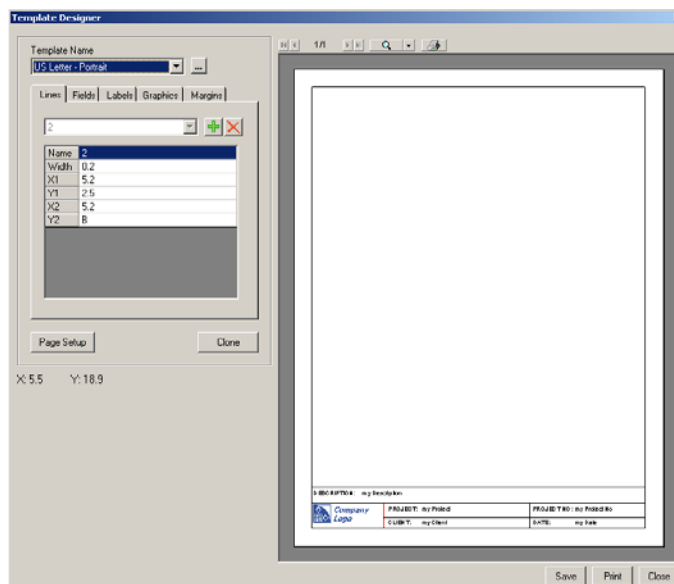
Enter “**R**” to have a line coordinate equal to the **right** margin location.

Enter “**T**” to have a line coordinate equal to the **top** margin location.


Enter “**B**” to have a line coordinate equal to the **bottom** margin location.

This is useful if you want the lines to automatically adjust as the page margins change. An example is shown below:

In this example, a line would be located at a height of 4 cm from the bottom of the page (Y1 and Y2 = 4) and the line would extend from the left page margin (X1 = L) to the right page margin (X2 = R) as shown below.



NOTE: To allow for quick recognition, the selected line, field or label is colored in **red** in the preview window. This will help you to quickly choose the starting and end points.



To remove existing lines, select the line then press the  button. The template designer settings will automatically be saved to the database; as such there is no **[Save]** button required in this dialogue.

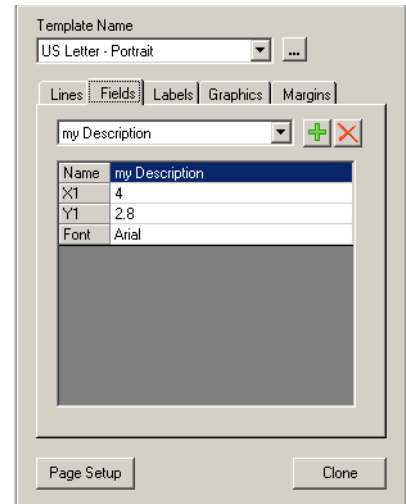
Creating Fields


The next two tabs in the template designer are **Fields** and **Labels**. These provide options for adding text to the template. The fields tab is used to create, position and label the Text fields. Beside each field, there will be a corresponding label.

The Text fields are the items which will show up when you load the template into the print preview window. They will be blank and prompt you to enter the specific details.

To create a new Field:

- Click  button and a new Field will be created with the name **new Field#**.
- Enter a unique name for the field.
- Enter the appropriate page coordinates for the field (the position refers to the page coordinate location for the bottom-left corner of the field). Remember to leave room for the Label which will typically be placed to the LEFT of the field.
- Select an appropriate font for the field. Click once in this field, and you should see a  button will appear. Press this button to access the font options for this field.



To modify an existing field, select this item from the combo box, and make the desired changes to the position, text value and/or font. The  button can be used to remove existing fields from the template.

The next step is to create a **Label** for the Text Field. As mentioned above, the Label is commonly positioned immediately to the left of the Field, using the same Y-coordinates. Therefore, take note of the Y-coordinates for the field(s) you just created, so they can be used for the corresponding label.


Creating Labels

The fixed text **Labels** are hardcoded into the template; this means that this item will always show up once the template has been selected, and will be Read-Only in the print options window. Beside each label, there is a **Field** entry, which receives the actual text entry for the printout when the template is loaded (information is entered into the fields in the print preview window).

For example, a label item would be **Date**; beside the date, there would be a **Date** field. When the template is loaded into the print options window, you will be prompted to enter a date in the **Date** field, which corresponds to the plot or report(s) which has been selected for printing.



AquaChem allows you to specify multiple labels at any location on the page. These labels generally contain static project information such as **Client**, **Project** number, and **Date**.


The **Labels** tab allows you to create, position and customize the Labels on the template.

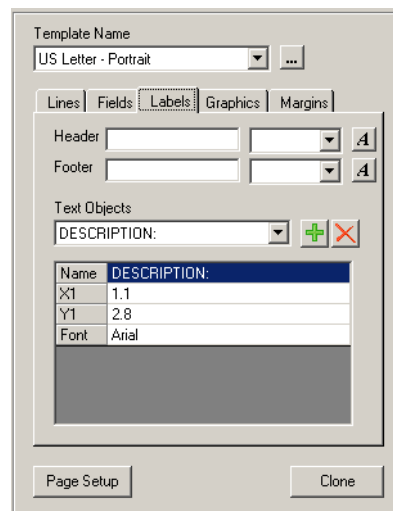
At the top of this dialogue, there are defaults for a **Header** and **Footer** for the template. The header position (**Left**, **Center**, or **Right**) can be adjusted along with the font (press the  button to access the font options for the Header or Footer). If you do not want a header or footer, simply leave this field blank.

In the lines below, you will see the list of labels.

To create a new label:

- Click  button and a new Label will be created with the name **new Text#**.
- Enter a unique name for the label.
- Enter the appropriate page coordinates for the label. (X1, Y1 refers to the page position for the bottom-left corner of the label). It is common to position the label to the left of the corresponding field. If necessary, use the preview window as a guide.
- Select an appropriate font for the label. Click once in this field, and you should see a  button appear. Press this button to access the font options for this label.

To modify an existing Label, select the item from the combo box, and make the desired changes to the position, text value and/or font. The  button can be used to remove existing Labels from the template.






Creating Graphics

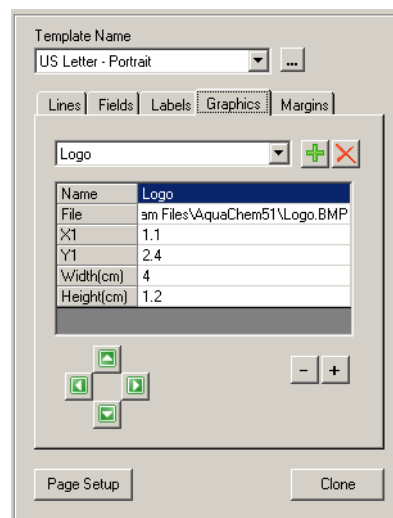
The **Graphics** tab in the template designer dialogue allows you to import graphics (.BMP, .GIF, or .JPG) into the template design.

You may set the position of the image on the page and specify the size as it will appear on the page. Each of the default Print Templates has been designed to accommodate an imported bitmap image of your company logo or your client's logo (or any image). AquaChem supports common graphics files .BMP, .GIF, and .JPG.

To import a new bitmap image file:

- Click  button and a new Graphics entry will be created with the name **new Picture#**.
- Enter a **Name** for the new graphics.
- Click once in the **File** field and a  button should appear. Click this button to browse the bitmap image file (filename.BMP) and then to import the image click **[Open]**.
- Enter the position coordinates of the bitmap (the position of the bitmap refers to the X-Y page coordinates for the top-left corner of the bitmap image).
- Enter the **Width** and **Height** of the selected bitmap file. The dimensions of the bitmap image refers to the Width and Height of the bitmap image as it will appear on the page. The bitmap image will be automatically re-sized (e.g. stretched or compressed) to fit the dimensions.
- You may also use the navigation and reduce/enlarge buttons to change the position and size of the label.

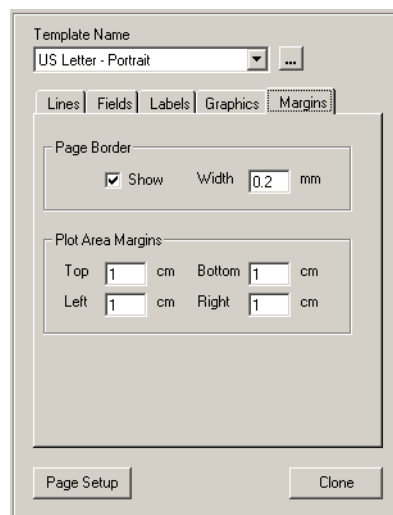
To modify an existing bitmap file, select the graphics from the combo box, and make the desired changes to the position, name, or dimensions. The  button can be used to remove existing bitmap logos from the list.



Adjusting Plot area Margins

The **Margins** tab in the template designer dialogue allows you to define the area that may be used for placing plots automatically. For example, if the template includes figure information at the bottom of the page, setting a margin of 5 or 6 cm will prevent the automatic plot placement from using this area and the plots will not overlap with these template elements. This is done by simply entering the distance (cm) from each side of the page you would like the display area to be confined to in the appropriate field (**Top**, **Bottom**, **Left**, and **Right**).

You can also turn the page border on or off using the **Show** check box and set the width of the page border line by entering an appropriate value (mm) in the **Width** field.



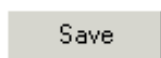
This concludes the steps for creating Print Templates.

3.1.11 Preferences

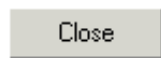
The **Preferences** dialogue loads the general user preferences and database options for AquaChem. These can be accessed only when the sample/station list is active. Several preferences and options are available and a detailed description for each of these are provided in the following section.

NOTE: If your database does not include some of the required parameters or if some of the required parameters are not properly mapped in the **Alias** section, then the applicable function(s) requiring these parameters will contain null values or erroneous information.

At the bottom of this dialogue, you will see two buttons:



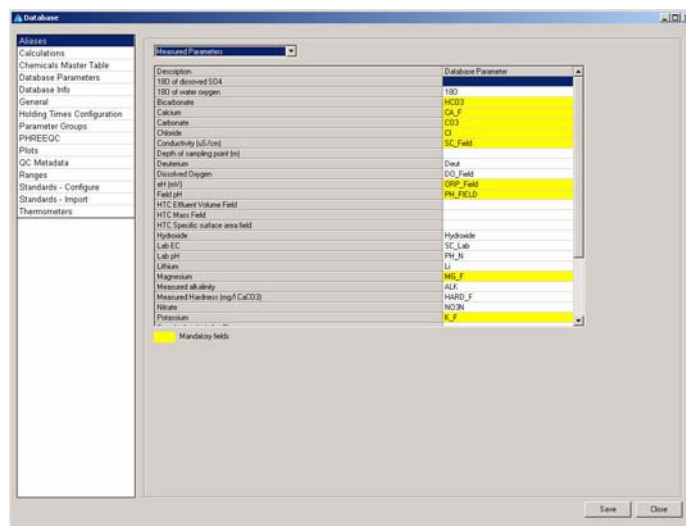
The **Save** button will save any changes you make to your database.



The **Close** button will close the dialogue, and return you to the main AquaChem window.

Aliases

This option allows you to adjust the parameters which are used for some of the internal calculations in AquaChem. Some parameter names have been hardwired to AquaChem for these calculations. For example, in order to calculate the hardness given by the sum of Ca and Mg, AquaChem must know if these parameters are called Ca or CALCIUM in the database.



Mandatory aliases which require to have an assigned parameter are highlighted in yellow. Aliases that have a white background may stay unmatched, however, certain functionalities may not be available. e.g. if the 18O and Deuterium aliases are not defined, the meteoric waterline plot and isotope related calculations will not be active, even if the respective parameters exist in the database.

For most purposes, it is recommended not to modify the original parameter names. However if you must rename a mandatory parameter, this section gives you the opportunity to map the database field to the parameter name in the database. An important alias is the “Measured Alkalinity”. Often carbonate species are measured in the field as alkalinity and expressed in mg/l CaCO₃. If this is the case, then the respective parameter must be matched to the Measured Alkalinity alias. After that, the electroneutrality calculation will be based on alkalinity rather than HCO₃ and CO₃ concentrations. Alkalinity may be plotted on Piper diagrams and PHREEQC will use this parameter when modeling the aqueous species distribution. After setting the measured alkalinity alias, make sure you return to the respective parameter and select the appropriate unit, as alkalinity units (mg/l CaCO₃, french degrees, german degrees, meq/L), differ from all other concentration units (mg/l, mmol/l etc.).

When you are dealing with very alkaline water samples and your data contains measured hydroxide concentrations, make sure that the “Hydroxide” alias is matched to the respective parameter since it is used for the alkalinity calculation.

Once you are finished in the **Aliases** tab, press **[Save]** to save any changes and **[Close]** to return to the main menu.

Calculations

The **Calculations** option allows you to set the equation coefficients and parameters that are used in the isotope and geothermal gradient calculations. It also provides information on the various calculations that are performed by AquaChem.

Isotopes

In the **Isotopes** frame, you may define coefficients for several functions for the isotopic calculations. With the exception of the meteoric water line, all equations have a highly regional character. If you want to make temperature or precipitation estimations, please find the respective coefficients for your study area. There are also parameters for ¹⁸O (Oxygen 18) and ²H (deuterium) equations: meteoric waterline, altitude-isotope and altitude-mean annual temperature relations. All of these functions are derived empirically. The coefficients for the isotope-altitude and the isotope-temperature functions are regional and should be used only in the area of calibration. Indicate the area of application of these functions in the field provided.

Geothermal Gradient

The **Geothermal Gradient** is used for a simple subsurface temperature estimation. These options allow you to specify which values are used to calculate the temperature at a given depth. The geothermal gradient is used in the geothermometer plot. It allows the user to estimate the subsurface temperature of waters produced by a borehole if the depth of the inflow zone is known. This temperature is then compared to geothermometer temperatures for the respective sample.

The formula for the Geothermal Gradient calculation is as follows:

The subsurface temperature $[T(z)] = \text{Average surface temperature} + \text{geothermal gradient} * \text{depth (km)}$.

The geothermal gradient may vary regionally; it is approximately 33 °C/km in most environments. Any changes made to the Geothermal gradient equation should be registered with AquaChem by pressing the **[Save]** button prior to closing the **Database** options dialogue.

Water Type (major ion definition)

First you may choose between an long and short name for water facies calculation. For the long version of the water type, the water type expression is calculated as follows:

- All concentrations are converted to meq/L.
- Values are then transformed from meq/L to meq%.
- All parameters which are below a certain limit (10% by default) are removed.
- The remaining ions are considered **Major ions**. These ions are ordered according to their percentage. Cations are ordered first, followed by the anions.
- The result is the **WaterType** string (e.g. Ca-Mg-HCO₃).

The threshold for determining if an ion is major is normally 10%; however in some countries 12.5% is more common. the higher the threshold, the smaller number of parameters that may exceed this threshold is. Usually, only major ions (Na, Ca, Mg, HCO₃, Cl, SO₄) will have a concentration that allows them to exceed the defined threshold. If you want to make sure, that other important elements such as potassium or nitrate appears in the water type expression, then you can achieve this by lowering the threshold value.

The second (short) way of calculating the water type consists of simply listing the most frequent cation and the most frequent anion, which results in one of the following combinations for most natural waters: Ca-HCO₃, Ca-Cl, Ca-SO₄, Na-Cl, Na-SO₄, Na-HCO₃.

The Redox Water Type option allows you to determine the water type based on the redox category of the sample. There are four redox categories (shown in the table below).

Category	Description	Description
A	strongly oxidized	contains oxygen, nitrate and sulfate, but no iron
B	weakly oxidized	contains nitrate and sulfate, but no oxygen or iron
C	weakly reduced	contains sulfate and iron, but no oxygen or nitrate
D	strongly reduced	contains iron and may contain hydrogen sulfide or methane, contains no oxygen or nitrate, sulfate reduced to some degree.
X	redox conflict	Sample contains a redox conflict, e.g., if hydrogen sulfide (strongly reduced) and oxygen (highly oxidized) are found in the same sample.

The Redox Category (water type) is determined by the following algorithm

The water type is displayed in the **Sample Details** window as a distinct parameter.

NOTE: The **Water Type** parameter cannot be edited; it is read-only. Also when changing the type of water type expression or when changing the threshold value for major ions, you must recalculate the water type expressions for all samples using the Tools>Calculators>Calculate Facies menu item.

Functions

The **Functions** frame lists all of the internal calculations performed by AquaChem, with reference information for each calculation provided in the fields below. All of the active functions (as indicated by a checkmark) will be available as database parameters which can be included in statistical comparisons and for plotting data. In this dialogue, you may select which calculations should appear in the function lists. For example, if you never use the enthalpy calculation, you may deactivate it here. In addition, you may define the default units for some of the functions.

The calculated values are displayed in the **Sample Details** window, under the **Calculated** tab:

The functions such as hardness, ion balance, etc. are hardwired to AquaChem; as such, it is not possible to change the parameters involved in these calculations. You may however, edit the name of the function or activate/deactivate a function. Deactivated functions will not appear in the list of functions within AquaChem. For functions that can be expressed in different units (e.g. hardness as °f, °g, mg/L CaCO₃) you may specify the default unit.

To edit the name of the function, press **View > Options** when the Sample Details window is active, and edit the options under the **Functions** tab.

AquaChem includes a number of common calculations for determining common geochemical parameters. Each of the available calculations (functions) is explained below.

The screenshot shows the 'Sample Details' window for 'Sample OW-2-93'. The 'Calculated' tab is selected, displaying a table of functions and their values. The table has three columns: Function, Unit, and Value. The functions listed include Electroneutrality, Sum of Anions, Sum of Cations, Sum of Ions, Calc TDS, Total Hardness, Carb. Hardness, Non Carb. Hardness, Calc Alkalinity, Calc Density, Calc TOC, and SAR. The values are displayed in the 'Value' column.

Function	Unit	Value
Electroneutrality	%	54.76
Sum of Anions	meq/l	18.15
Sum of Cations	meq/l	62.08
Sum of Ions	meq/l	80.22
Calc TDS	mg/l	2049.25
Total Hardness	mg/l CaCO ₃	2591.00
Carb. Hardness	mg/l CaCO ₃	124.96
Non Carb. Hardness	mg/l CaCO ₃	0.00
Calc Alkalinity	mg/l CaCO ₃	2.50
Calc Density	g/cm ³	1.002
Calc TOC	mg/l	0.07
SAR		2.01

Calculated Alkalinity

Acid neutralizing capacity. Generally equal to the concentration of $2\text{CO}_3^{2-} + \text{HCO}_3^- + \text{OH}^- - \text{H}^+$.

Calculated Density

Calculated density based on the partial molal ionic volumes. This value is calculated as follows:

$$D = \frac{\left[1000 - v + \frac{TDS}{1000} \right]}{1000}$$

where

v is the molar volume: $v = \sum(n_i \cdot v_i)$,

n_i is the concentration in mmol/l of ion i and,

v_i is the partial molar volume at 25°C

Partial molal ionic volume coefficients are from Owen and Brinkley, 1941 and for more details, see Hounslow, 1995, p. 58

Calculated TDS (Total Dissolved Solids)

TDS is a measure of the evaporation residue at a given temperature. It can also be thought of as the mass of all ions. It is important to note that the calculated TDS is only an approximation and it is always better to have a measured value. TDS is calculated as follows:

First the sum in mg/L of $\text{Na} + \text{K} + \text{Mg} + \text{Ca} + \text{Cl} + \text{SO}_4$ is calculated. Then AquaChem checks if $(\text{Ca} + \text{Mg} - \text{SO}_4) > (\text{HCO}_3 + \text{CO}_3)$. This is done because TDS is not simply the sum of ions but the weight of residue you would expect if you evaporate the entire sample. In waters that contain an excess carbon $(\text{Ca} + \text{Mg} - \text{SO}_4) < (\text{HCO}_3 + \text{CO}_3)$ you may not add all carbon to the residue since bicarbonate will partly outgas as CO_2 during evaporation. Only the quantity of HCO_3 which can be precipitated as calcite and dolomite is added to the estimate of the TDS. SO_4 is deduced from $\text{Ca} + \text{Mg}$ because gypsum is built first and the Ca that is used up by this process cannot be used any more to build calcite. If $(\text{Ca} + \text{Mg} - \text{SO}_4) > (\text{HCO}_3 + \text{CO}_3)$ then all HCO_3 and CO_3 can be added to the TDS because there is enough $\text{Ca} + \text{Mg}$ to build calcite and dolomite.

Total Hardness

The sum of ions that can precipitate from water as calcite or dolomite. Generally, the sum of Ca and Mg, expressed in meq/L or mg CaCO₃/L, or in degrees.

$$100 \text{ mg CaCO}_3/\text{L} = 1 \text{ mmol Ca}^{2+}/\text{L} = 2 \text{ meq Ca}^{2+}/\text{L}$$

The Degrees equivalents are as follows:

$$1 \text{ German Degree} = 17.8 \text{ mg CaCO}_3/\text{l}$$

$$1 \text{ French Degree} = 10 \text{ mg CaCO}_3/\text{l}$$

Non Carbonate (Permanent) Hardness:

Parts of Ca and Mg in excess of HCO₃.

Carbonate (Temporary) Hardness:

Parts of Ca and Mg which are balanced by HCO₃ and thus can precipitate as CaCO₃.

Residual Sodium Carbonate (RSC)

The RSC has the following equation: $\text{RSC} = (\text{CO}_3^- + \text{HCO}_3^-) - (\text{Ca}^{2+} + \text{Mg}^{+2})$

It is similar to the SAR in that it expresses the sodium content in relation with Mg and Ca. This value may appear in some water quality reports although it is not frequently used. If the RSC < 1.25 the water is considered safe If the RSC > 2.5 the water is not appropriate for irrigation.

Langelier Saturation Index (LSI)

The Langelier index is a popular way of expressing the equilibrium state of a solution in respect to calcite.

- Index between -0.5 and +0.5: the water is balanced
- Index < -0.5: the water is Corrosive.
- Index > +0.5: the water is scale forming.

$$\text{LSI} = \text{pH} - \text{pHs}$$

where pH is the measured pH and pHs is the pH where the water sample would be in equilibrium. The pHs is estimated from temperature, alkalinity, hardness and total dissolved values.

$$\text{pHs} = (9.3 + a + b) - (c + d)$$

$$a = \log(\text{TDS}) - 1) / 10$$

$$b = -13.12 * \log(\text{temp} + 273.15) + 34.55$$

$$c = \log_{10}(\text{Ca as CaCO}_3) - 0.4$$

$$d = \log(\text{Alkalinity})$$

Note that the LSI is only an approximation of the real saturation state. Whenever possible you should use the PHREEQC modeled saturation index of calcite or dolomite in your studies.

Ryznar Stability Index (RSI)

The Ryznar stability index (RSI) attempts to correlate an empirical database of scale thickness observed in municipal water systems to the water chemistry. Like the LSI, the RSI has its basis in the concept of saturation level. Ryznar attempted to quantify the relationship between calcium carbonate saturation state and scale formation. The Ryznar index takes the form:

$$\text{RSI} = 2(\text{pH}_s) - \text{pH}$$

The empirical correlation of the Ryznar stability index can be summarized as follows:

- RSI << 6 the scale tendency increases as the index decreases.
- RSI >> 7 the calcium carbonate formation probably does not lead to a protective corrosion inhibitor film.
- **RSI >> 8** mild steel corrosion becomes an increasing problem.

ElectroNeutrality

(Cations - Anions)/(Cations + Anions), expressed as a percent. The Electro Neutrality is a common quality control method.

Sum of Anions:

Sum of all measured anions for the sample, in meq/l.

Sum of Cations:

Sum of all measured cations for the sample, in meq/l.

Sum Ions:

Sum of all ions in your sample. This can be expressed in mmol/l, meq/l, mg/L.

Exceeds_Lev1, Exceeds_Lev2, Exceeds_Lev3

The functions “Exceeds_Lev1, Lev2, Lev3” returns “-1” for samples which contain at least one concentration exceeding the specified water quality standard and they return 0 if no concentration has been exceeded. It is a convenient way of finding samples which should be verified and analyzed more carefully. This is especially effective if you are dealing with many different parameters that might exceed the guidelines.

SAR (Sodium Adsorption Ratio)

The sodium adsorption ratio is an important water quality criterion for irrigation waters. It is used in the Wilcox plot.

$$SAR = \frac{Na}{\left(\frac{Ca + Mg}{2}\right)^{0.5}}$$

NOTE: The concentrations used in the formula are in meq/L.

MH (Magnesium Hazard)

Magnesium is considered to be harmful for plants, but the effect is reduced by the presence of calcium. Magnesium Hazard was proposed by Szabolcs and Darab (1964).

$$MH = \frac{Mg}{(Ca + Mg)} \times 100$$

Units are in meq/l. MH > 50 is considered to be harmful for irrigation water.

NOTE: The Concentrations used in the formula are in meq/L.

Calculated TOC

This function calculates the total carbon within all organic species. Carbon from inorganic species such as HCO₃, CO₃, CO₂ is not included.

Calculated TOX

Calculated total organic halogens. Sum of all halogens (e.g. Cl⁻, Br⁻, I⁻, etc.) present in ug/L, mg/L or g/L in all organic parameters (thus, Cl from Cl⁻ or CL2 is excluded).

z(18O)

Average infiltration height as a function of oxygen 18 isotopic composition. Coefficients are valid only for a very limited zone, and must be established for your area of interest using empirical data.

T(18O)

Average temperature of infiltration zone as a function of oxygen 18, isotopic composition. Coefficients are valid only for a very limited zone, and must be established using empirical data.

T(2H)

Average temperature of infiltration zone as a function of deuterium. Coefficients are valid only for a very limited zone.

z(2H)

Average infiltration height as a function of deuterium. Coefficients are valid only for a very limited zone.

Temperature (Z)

Estimated Temperature for a given depth and geothermal gradient.

Temp > H H2O (l)

Enthalpy of liquid water as a function of temperature (t). This is a polynomial function:

$\text{Temp} = \text{enth}(\text{water})$

$\text{value!} = a(1) + a(2) * t + a(3) * t^2 + a(4) * t^3 + a(5) * t^4 + a(6) * t^5 + a(7) * t^{-1} + a(8) * t^{-2} + a(9) * \log_{10}(t)$

where

$$a(1) = 418.84$$

$$a(2) = 10.286$$

$$a(3) = -0.05092$$

$$a(4) = 0.00026309$$

$$a(5) = -0.00000069303$$

$$a(6) = 0.00000000074566$$

$$a(7) = -1209.8$$

$$a(8) = 11.99$$

$$a(9) = -353.76$$

For more details please refer: Fournier & Potter, 1972.

Temp > H H2O (v)

Enthalpy of water vapor as a function of temperature. This is a polynomial function:

$$\text{Temp} = \text{enth}(\text{water vapor})$$

$$\text{enthalpy} = a(1) + a(2) * t + a(3) * t^2 + a(4) * t^3 + a(5) * t^4 + a(6) * t^5 + a(7) * t^{-1} + a(8) * t^{-2} + a(9) * \log_{10}(t)$$

where

$$a(1) = 2035$$

$$a(2) = -5.0499$$

$$a(3) = 0.057399$$

$$a(4) = -0.00030426$$

$$a(5) = 0.00000079095$$

$$a(6) = -0.00000000086968$$

$$a(7) = 1342.4$$

$$a(8) = -13.298$$

$$a(9) = 396.29$$

For more details please refer: Fournier & Potter, 1972:

SiO2 > H H2O (l)

Enthalpy of liquid water as a function of dissolved silica. This is a polynomial function:

$$\text{enth} = a(1) + a(2) * \text{SiO}_2 + a(3) * \text{SiO}_2^2 + a(4) * \text{SiO}_2^3 + a(5) * \log_{10}(\text{SiO}_2)$$

where

$$a(1) = -42.198$$

$$a(2) = 0.28831$$

$$a(3) = -0.00036686$$

$$a(4) = 0.00000031665$$

$$a(5) = 77.034$$

For more details please refer: Fournier & Potter, 1972:

SiO2 > H H2O (v)

Enthalpy of water vapor as a function of dissolved silica. This is a polynomial function:

$$\text{enth} = a(1) + a(2) * \text{SiO2} + a(3) * \text{SiO2}^2 + a(4) * \text{SiO2}^3 + a(5) * \log_{10}(\text{SiO2})$$

where

$$a(1) = -3.5532$$

$$a(2) = 0.146$$

$$a(3) = -0.0004927$$

$$a(4) = 0.0000012305$$

$$a(5) = -0.00000000049421$$

For more details please refer: Fournier & Potter, 1972:

Exceeds_Lev1, Exceeds_Lev3, Exceeds_Lev3

These functions return true (-1) for all samples, where at least one parameter exceeds the respective standard level (1 2 or 3). This function is convenient when used in a query expression. If you query for exceeds_lev1 = -1 then all samples having values exceeding the first level of the current water standard will be highlighted in the sample list.

Exceeds_Solubility

Returns “true” (-1) if solubility is exceeded and the solution is includes a chemical which solution is above the specified solubility.

Exceeds_NO

Returns “true” (-1) if natural occurrence is exceeded by one of the measured concentrations. The natural occurrence range can be defined in the database screen on the range tab.

Season

Returns the season number for the sample (i.e. 1=Spring, 2=Summer, 3=Fall, 4=Winter - based on Northern Hemisphere seasons). Note that this is a convenient way to find and select all samples that were sampled in the same season but in different years.

- Spring: 21/3 - 20/6
- Summer 21/6 - 20/9
- Fall 21/9 - 20/12
- Winter 21/12 - 20/3

On the southern hemisphere as defined in the preferences screen: 1=Fall, 2 equal Winter, 3=spring, 4=Summer

Month

Returns the month for the sample (i.e. 1=January, 2=February, etc.). Again, using this function in the find screen might be convenient to select data for calculating month based statistics over several years.

Year

Returns the year for the sample.

Once you are finished in the **Calculations** section, press **[Save]** to save the changes and **[Close]** to return to the main menu.

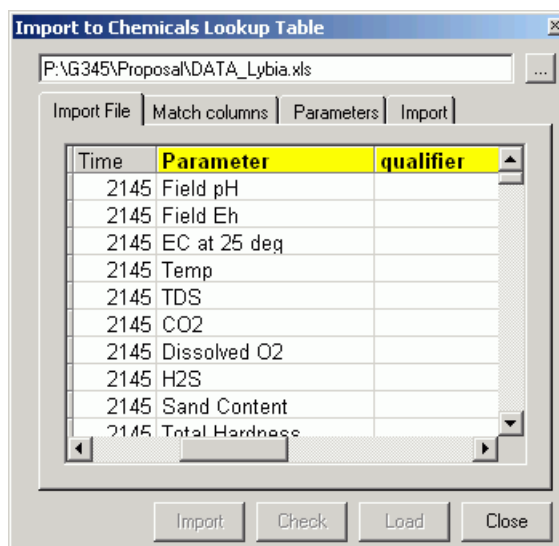
The calculated parameters are treated by AquaChem as regular database parameters with respect to plotting, searches, or statistical calculations. However, in order for the built-in calculations to work, their corresponding database parameters must be included in the database. For example, the database must include Ca and Mg in order to calculate hardness, and both of these parameters must be identified with an Internal Name of 'Ca' and 'Mg' respectively.


Chemicals Master Table

This option allows you to maintain the chemicals master table, which is stored within the Aquachem database file. This table is an extensive list of predefined chemicals, which may be used as templates to create new parameters in the AquaChem project files. You may manually add new parameters, delete parameters, or change fields in existing parameters.

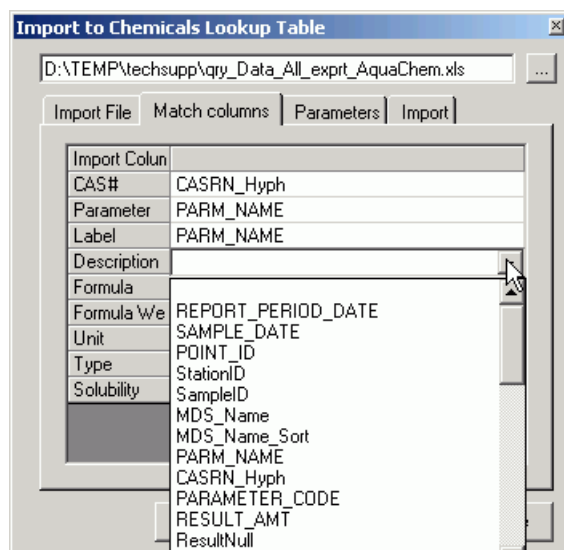
Importing Parameters

The import file typically contains dozens of chemicals that need to be created manually in your database before the respective values can be imported. Often these files have the “one value per row” format as shown below. In this format it is very tedious to extract individual parameter names. The **[Import]** button allows you to automate the extraction of these parameters and adds them to the chemicals master table and/or the current database parameters. After pressing the **[Import]** button you will see the following screen:

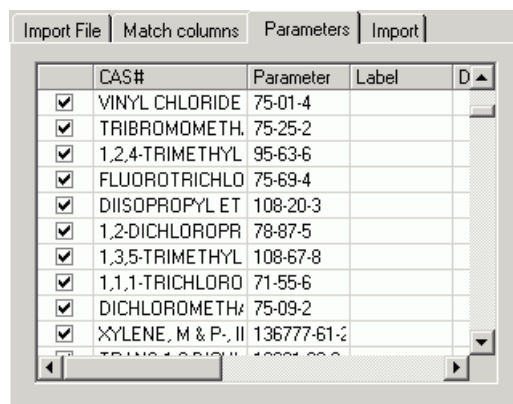


First, you need to specify the import file by pressing the  button. The grid below will be filled with the file contents.

On the **Match columns** tab you have to match the import file columns to the respective column in the chemicals master table. The CAS number and the parameter name are mandatory.

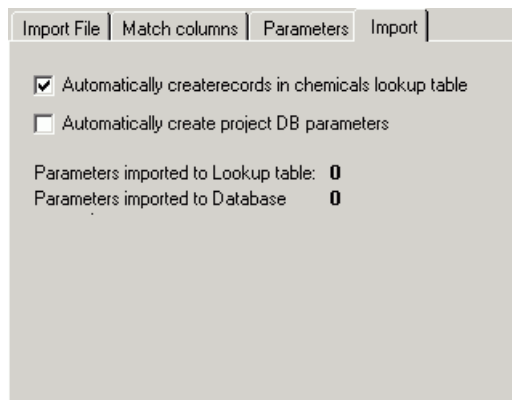


Press the **[Load]** button to proceed. This will extract all distinct parameter names found in the specified column and switches the display to the parameters tab. The **Parameters** tab should now contain the list of extracted parameters as shown below. Each record is preceded by a checkbox, indicating whether this parameter will be imported in the following import procedure.



Press the **[Check]** button in order to test the parameters for entries having different names but the same registry number. This may be due to a typo in the parameter name, in which case the parameter should be discarded. This can occur as well because of having the same parameter measured in two different ways, e.g. filtered and unfiltered. You might want to add such parameters to the database parameters, but not to the Chemicals Master Tables, which should contain one record per CAS registry number only. The system prompts you when encountering duplicate registry numbers and

unchecks the records, if you decide to discard it. Once the parameter check is ended, the **Import** tab is shown.



This final tab lets you specify if the parameters should be imported to the Chemicals Master Table only or in parallel to the current project database parameters as well. We recommend to import parameters to the Chemicals Master Table only. Then review and complete the parameters with missing formula weight and formula information. Finally, go back to the **Parameters** and add new parameters from the updated Chemicals Master Table.

Press the **[Import]** button to start the import procedure. Note that you will have to confirm the import of parameters where an identical CAS registry number already exists in the target table. If a parameter is to be imported having the same name as a parameter already existing in the target table, this parameter will be simply ignored. At the end of the import process the number of successfully created parameters in the target table will be displayed on the **Import** tab.

Database Parameters

The **Database Parameters** option allows you to modify the data structure of the current AquaChem database project by adding or deleting parameter fields from each of the four parameter categories. This option can also be used to modify the attributes of each parameter (e.g. Display label, Internal name, Molecular weight, Charge, Units, etc.) or to change to order in which the parameters are displayed in the Sample Details window.

In an AquaChem database there are several categories of parameters:

- **Station Description Parameters**
- **Sample Description Parameters**
- **Measured Parameters**
- **Analysis Metadata**, and
- **Modeled Parameters**

Mandatory parameters are highlighted in yellow. A detailed description of each parameter category follows below:

Station Description Parameters

These are parameters which contain general information about the station. This data is entered once for every individual station. The station can then be assigned to a single sample or multiple samples. Typical parameters for the stations are: station name, ID, location, area, X,Y coordinates, elevation, and well depth. Station Description parameter values can be text or numbers. An AquaChem database may contain a maximum of 255 Station Description parameters.

Sample Description Parameters

These are parameters which contain general information about the sample. This category typically includes alphanumeric or data types such as sample ID, sampling date, sampling personnel, etc. Sample Description parameter values can be text or numbers. An AquaChem database may contain a maximum of 255 Sample Description parameters.

Measured Parameters

These are parameters that are either measured at a sampling location (in the field) or derived from a laboratory analysis of a water sample taken from that location. These parameters will contain numerical values for your sample (ex. concentrations for cations, anions, organic contaminants, pH, TDS, etc.). For these parameters, you may enter a comment for every analyzed value as well as an acceptable Range. An AquaChem database may contain unlimited number of Measured parameters.

Analysis metadata

In AquaChem, each measurement may be further described with meta data fields. The initial database includes the most commonly used fields: Method detection limit, Protocol (Method), outlier flag, solubility, and comment. Depending on your field of activity you might want to add meta data fields such as the Practical Quantification Limit (PQL) or the information whether this element has been filtered after sampling. AquaChem allows you to add an unlimited number of meta data fields to the measurement table. Meta data information is only visible in the **Sample Detail** screen where each parameter appears as an additional column behind the measured value. Note that you have to set the respective to Visible using the **Sample Detail Option** screen.

Modeled Parameters

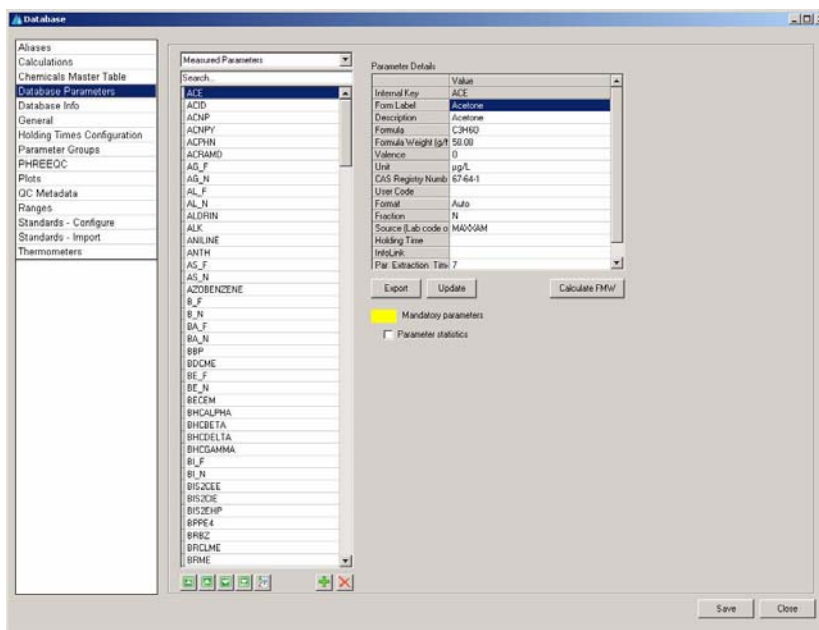
AquaChem provides an integrated graphical interface for geochemical modeling using PHREEQC where the data for each sample in the database can be used as input data for the PHREEQC simulations. Modeled Parameter values are calculated by PHREEQC during a simulation and read back to the database.

The Modeled Parameters are very similar to numerical parameters, however you may not add comments, description, or define Ranges. Modeled Parameters appear in a separate tab in the Sample Details window.

For more details on the Modeled Parameters and PHREEQC modeling, please refer to Chapter 6: Tools, and Chapter 7: Geochemical Modeling with PHREEQC (Basic).

Parameter Details

For each Parameter category, you will see a list of parameters belonging to that category appearing on the left side; on the right side, you will see the description and details for each parameter, as shown below:



At the bottom of the Parameters list, you will see several buttons. The function of each button (in order from left to right) is as follows:



Move to top: moves a parameter to the top of the list



Move up: move the parameter up



Move down: move the parameter down



Move to bottom: move the parameter to the bottom of the list



Sort alphabetically: sorts parameter list alphabetically



Add new parameter: adds a new parameter



Delete parameter: deletes the selected parameter(s)

Each parameter has a corresponding name, label, and info, so that it can be easily identified in the database, and incorporated in to plots, calculations, and modelling with PHREEQC. Below is a list of the required fields for the various parameter types.

Sample Description Parameters and **Station Description Parameters** require the following **Parameter Details**:

- Internal Key
- Form Label
- Description
- Type

One of the new features of AquaChem is the ability to modify the type of the existing parameters for the station and the sample table. Depending on the **Type** of a parameter, different details need to be provided:

Type **Integer, Long, Single, Double**:

- Unit
- Format

Type **Text**:

- Length
- List of Values (optional)

Type **Memo**

- Length

Type **Boolean** and **Date**:

- No additional parameter details

A **Measured Parameter** requires the following **Parameter Details**:

- Internal Key
- Form Label
- Description
- Formula
- Formula Weight (g/Mol)
- Solubility
- Valence
- CAS Registry Number
- Unit
- Format




A **Modeled Parameter** requires the following **Parameter Details**:

- Internal Key
- Form Label
- Description
- Formula
- Formula Weight (g/Mol)
- Valence
- Unit
- Format

NOTE: It is not necessary to have values entered for each field for each parameter. However for reports and plots, it is recommended that you enter as much information as possible.

The following is a brief summary of each of these **Parameter Details**:

Internal Key	This is the internal name defined when the parameter is first created; once a parameter has been created, the internal key cannot be changed.
Form Label	Display label assigned to the parameter so that it can be distinguished in the Sample/Station Details window, and other AquaChem windows and dialogues.
Description	Brief description of the parameter. This information is entered only in this location, and is designed to provide the user with a quick reference. (This field may remain blank if desired).
Length	Set the character length of the field
Formula	Chemical formula for the parameter.

List of Values	Allows you to create a unique pick list for this parameter. To edit the list of values, click the  button in the List of Values field. When editing the respective parameter in the sample or station window, only the predefined values can be selected from a combo box.
Formula Weight	Formula weight for the measured parameter. For many measured parameters, there is an option to calculate the formula weight (select a measured parameter then you will see a Calculate FMW button appearing below this dialogue. Press this button and AquaChem will automatically calculate the weight of the parameter, provided the formula is entered and available in the database).
Solubility	Set the solubility for the measured parameter.
Valence	Positive or negative charge (if applicable).
CAS Registry Number	For measured parameters, if you know the CAS (Chemical Abstracts Service) Registry number, you can enter it in this field. Otherwise, you may leave this field blank.
Unit	Allows you to select the units that will be applied for the selected parameter (mg/L, meq/L, etc.). For all other parameters without a formula, an editable list of units is displayed. To access this list, simply click in the field beside Unit and press the  button to load the list of available units. Note that when changing the unit for a parameter and the database already contains data, existing values are not changed automatically. This change has to be done manually using the Edit > Replace option after saving the database configuration. E.g. when changing the unit for copper from mg/L to ug/l, all copper values have to be multiplied by a factor of 1000.
Format	Displays the format for a numerical parameter. This field determines how many decimal places will appear for a numerical parameter. To change the display format, simply click in the field beside Format and press the  button to load the Format options dialogue. You can then increase/decrease the number of Decimal places using the up/down buttons or select the Auto or Scientific format for the parameter.
Type	Set the parameter data type. There are nine data types in the AquaChem database: <ul style="list-style-type: none"> • Boolean • Integer • Long • Single • Double


- Date
- Text
- Memo

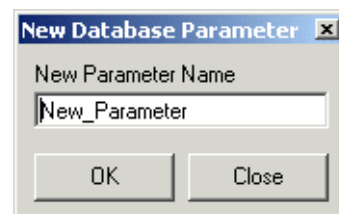
The numeric parameter fields do not accept text values such as "ND" or "NA" and will return a value of zero for any such text entry. However, AquaChem does accept "greater than" and "less than" values (e.g. <0.01). In addition, AquaChem allows you to recognize or ignore the "greater than" and "less than" values on graphs (see the **File > Preference**, and browse to the **Censored Data** tab to edit these options).

The next section will describe how to add new parameters to your database, and create new parameters.

Adding/Creating New Parameters


If you are adding a **Station Description** or **Sample Description** parameter to your database, follow the steps below:

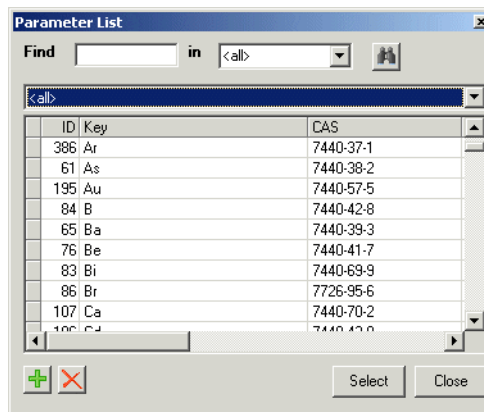
- Select **File > Database** from the main menu (if you have not already done so) and go to the **Parameters** tab.
- Select the **Station Description** or **Sample Description** parameter category from the combo box in the upper left corner of the dialogue.
- Press the  button and a **New Database Parameter** dialogue will appear (as shown on the right-hand side).
- In this dialogue type in the name of your new parameter. This name will be used as the Internal Key, Form Label, and Description for the parameter.
- Click **[OK]** and this will return you to the list of parameters.
- Proceed to fill in the required **Parameter Details** for this parameter.



If you are adding a **Measured Parameter**, then follow the steps below:

- Select the **Measured Parameters** category from the combo box in the upper left corner of the dialogue.


- Press the  button and you will see a **Parameter List** dialogue (as shown on the right-hand side).
- Choose a parameter from the list of available parameters; to add multiple parameters, press the **<Ctrl>** key (on your keyboard) while you select multiple parameters from the source list. You may use the **Find** option to run a query for a parameter name to see if it is available in the AquaChem chemical database.

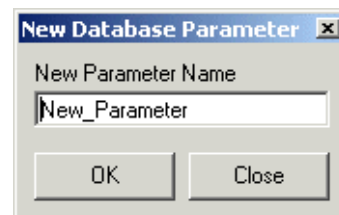


Use the combo box above the parameters list to choose from the various measured parameter categories available in AquaChem (Inorganic, Organic, VOC's, etc.).


- Press the **[Select]** button and the parameter(s) will be added to your database.

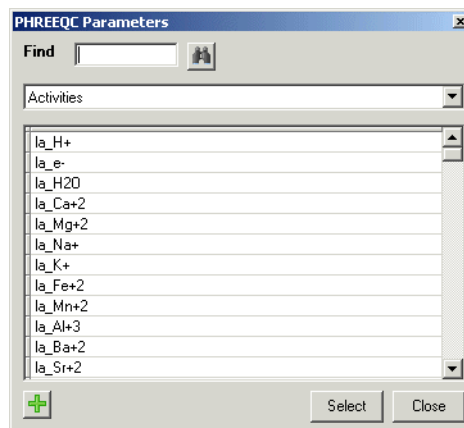
If the parameter that you want to add is **NOT** available in this list, then:

- Click the  button (in the lower-left corner of the dialogue) and a **New Database Parameter** dialogue (as shown to the right) will appear.
- Type in the name of your new parameter (this may be a formula or full chemical name). This name will be used as the Internal Key, Form Label, and Description for the new parameter.
- Click **[OK]** and you will return to the **Database** options dialogue.
- Proceed to fill in the required **Parameter Details** for this parameter.



If you are adding a **Modeled Parameter**, follow the steps below:


- Select the **Modeled Parameters** category from the combo box in the upper-left corner of the dialogue.
- Press the  button and you will see a **PHREEQC Parameters** dialogue (as shown on the right-hand side).
- Choose a parameter from the list of available parameters. To add multiple parameters, press the **<Ctrl>** key (on your keyboard) while you select multiple parameters from the source list. You may use the **Find** option to run a

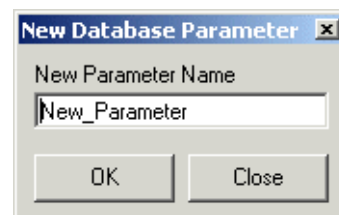


query for a parameter name to see if it is available in the AquaChem chemical database. Use the combo box at the top of this dialogue to choose from the various parameter categories available in AquaChem (**Activities**, **Saturation Indices**).

- Press the **[Select]** button and the parameter(s) will be added to your template.


If the parameter that you want to add is **NOT** available in this list, then:

- Click the  button (in the lower-left corner of the dialogue) and a **New Database Parameter** dialogue (as shown to the right) will appear.
- Type in the name of your new parameter. It is important that the name for new modeled parameter matches the names used in the PHREEQC thermodynamic database for phases (minerals) or species. When selected from the list, this will ensure that the names are matched successfully. This name will be used as the Internal Key, Form Label, and Description for the new parameter.
- Click **[OK]** and you will return to the **Database** options dialogue.
- Proceed to fill in the required **Parameter Details** for this parameter.

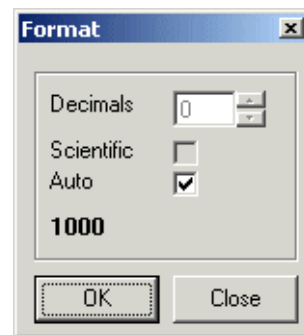


NOTE: Parameter names should contain numbers and letters only; do not use any other characters, and/or spaces in the parameter names.

Upon returning to the Parameter list, you must then enter the remaining required parameter details including the description, units, formula, format etc. All measured chemical parameters should have a formula weight specified. If you do not know the molecular weight of a chemical parameter, press the **Calculate FMW** button below the parameter details and AquaChem will read the formula from the field and insert an appropriate molecular weight according to the Periodic Table of Elements. Cations and Anions must be accompanied by their **Valence** (positive or negative charge).


The Format field is set to **Auto** by default. To choose a new display format for a parameter, click once in the **Format** field, and then press the  button and you will see the dialogue as shown on the right-hand side.

The **Format** options allow you to specify the number of **Decimal** places for measured parameters, and specify if the value will be in **Scientific** or whole number format. If **Auto** is selected, then AquaChem will determine the optimal number of decimal places for that parameter. This may result in a “hiding” of a 0 after a decimal place (for example, 7.10 may be displayed as 7.1). To correct



this, disable the Auto option, and increase the number of decimals to the desired amount. Once you are finished, press **[OK]** to accept changes or **[Close]** to return to the previous dialogue.

Deleting Parameters

To remove a parameter from your database, simply select one or several parameters from the parameters list on the left side of the dialogue and click the  button.

Exporting Parameters

On various occasions, it will be useful to save all parameters and respective settings to a file for documentation or review purpose. This may be accomplished by pressing the Export button below the parameter details grid. The information may be saved to in text or Excel format.

Update Parameters

The central Aquachem.mdb database holds a master table including hundreds of organic and inorganic chemicals. This is the list you may choose from when adding new parameters to the database. If a parameter has been created manually and the entry misses the formula or formula weight, you may use the **Update** button to copy the missing information from the chemicals master table to your database parameter. Note that for this the CAS registry number must be filled, since it is internally used for matching parameters. Also, if the parameter does not exist in the Master chemical table, the update process will not return any results.

Mandatory Parameters

AquaChem has several parameters which are required for EVERY database, and therefore cannot be deleted from the database. These parameters are needed by various plots, reports, and calculations. Mandatory parameters are highlighted in yellow and when selected, the **[Delete]** button becomes disabled.

The following is a list of mandatory parameters for an AquaChem database.

Mandatory Sample Description Parameters:

- Sample_Date
- SampleID
- WaterType

Mandatory Station Description Parameters:

- StationID

- X
- Y

Mandatory Measured Parameters:

Ca	HCO3	SAMPLE_DEPTH
Cl	K	Si
CO3	Mg	SO4
COND	Na	TDS
DENSITY	pH	TEMP
Eh		

Once you have made changes to the Parameters in your database structure, press **[Save]** to accept these changes.

Parameter Statistics

The Parameter Statistics frame allows you to view parameter statistics, e.g., Min, Max, Avg and Count, while browsing the parameters in the parameters list.

Database Info

This options gives you some basic information about your database including the template it is based on and size. As well as when it was created, modified and last accessed.



The **Compress** button will compress your database. Your project database file size may become large if you have a large number of samples, and are continuously deleting samples. Compressing the database will compact and repair your database, and may make the file size more manageable. AquaChem will create a backup copy of the uncompressed database file, with the file extension .SAV. Should you encounter problems with the new compressed file, simply delete this, and restore the previous file by renaming the .SAV file with the extension .AQC.

The AquaChem project database files are saved in a Microsoft Access 2000 relational database format. The projects are saved with a unique file extension (.AQC). This is done for two reasons:

1. This prevents any confusion or possible corruption of the file if it were to be accidentally opened in MS Access, and critical changes were made to the data structure or even to the data itself; and

2. This ensures that only AquaChem projects are opened in AquaChem, and that non-supported files that (that could potentially result in corruption or data loss) are not opened.


However, it is still possible to view/edit an AquaChem database file using MS Access. Simply launch MS Access and open this database file, using the **File > Open** command.

NOTE: It is recommended that you first make a back-up of your database file, before modifying it in the MSAccess environment.

An AquaChem database may contain an unlimited number of samples, depending on the number of parameters it uses and the amount of memory available on your PC. Working with too many samples can slow some operations even though you are only working with a portion of the database. Therefore, working with several databases containing less than 5,000 samples may be more efficient than using one large database.

To view and modify the AquaChem database structure, select **File > Preferences** from the main menu while the Samples/Stations list is active.

General

In the **Interface Colors** dialogue, you can specify the AquaChem main program window **Background** color. Simply click on the  button and a standard windows **Color** options dialogue will load allowing you to choose another color.

Use the **Project Location** combo box to select the hemisphere for your project site. Depending on the hemisphere, the seasons may be reversed. The seasons are counted beginning with Spring (March 21 to June 20), in the Northern Hemisphere and Fall in the Southern Hemisphere.

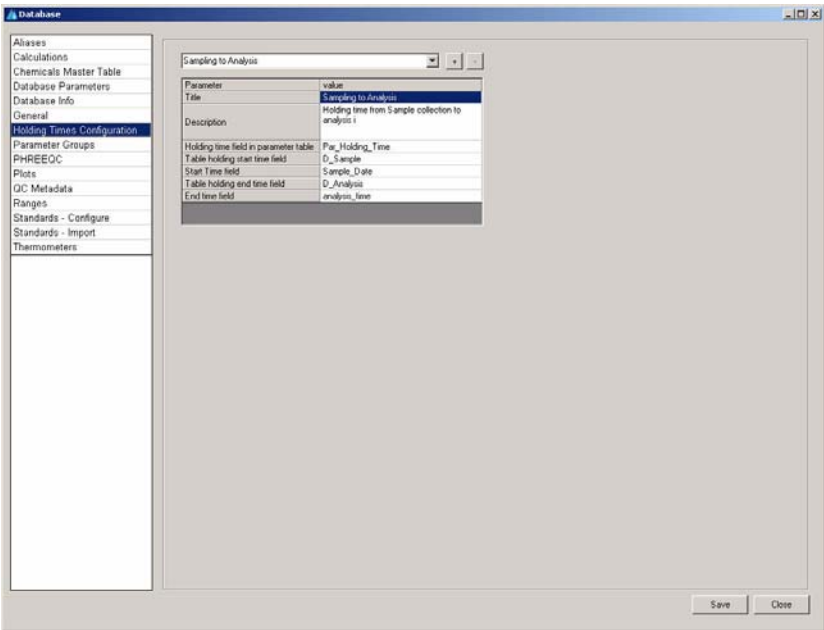
To view which season returns which number, consult the **Calculations** tab of the **Database**.

Use the **Document Root** field to specify the path to the folder, containing documents that are referenced from AquaChem records. Sample or station text fields may contain filenames or internet addresses. For filenames, you may indicate the full filename including the path or only the filename. In latter case, this file must be reside in the documents root folder. This is convenient, if the documents have to be moved to a different location. Since all reference files reside in the root folder, only the document_root parameter has to be changed to indicate the new location. If the files are specified including the full file path, then all records referencing the moved files must be updated.

Additionally you have an option to automatically open the last AquaChem project when you Startup the program.

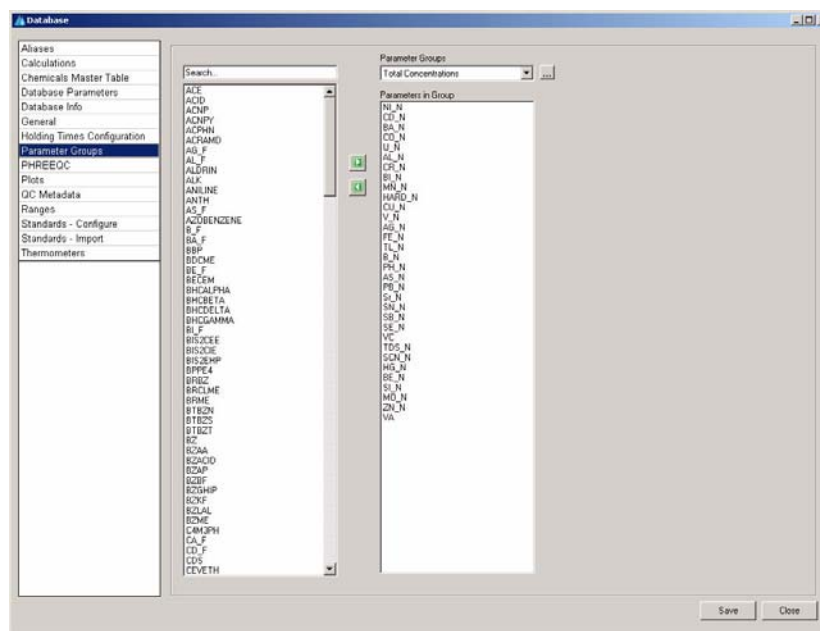
Holding Times Configuration

This option allows you to configure the tables and fields used to run a holding times analysis.





Parameter Groups

The **Parameter Groups** options allow you to sort measured numerical parameters into groups, providing a quick and easy view of specific sample data accessible in the Sample Details window.



For your convenience, AquaChem comes with the following pre-defined parameter groups.







- Show analyzed values only (hides all parameters for which there are no assigned measurement values)
- Anions (e.g. Cl^- , F^- , SO_4^{2-} etc.)
- Cations (e.g. Ca^{2+} , Na^+ , K^+ , Mg^{2+} , etc.),
- Gas (O_2 , CO_2 , NO_2 , etc.)
- Isotopes (^{18}O , ^2H , ^{14}C , etc.)
- Organic Chemicals (BTEX, Vinyl Chloride, etc.)
- Overview (pH, TDS, Cond, Major ions)

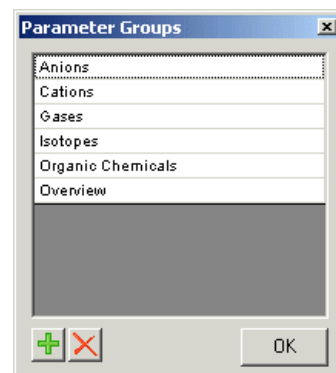
By selecting a parameter group from the combo box, you can view which parameters belong in a selected group; one parameter may belong to multiple parameter groups. To add/remove parameters from a parameter group, simply select the parameter from either the full list on the left side, or the group list on the right side, and press the  or  button to move the parameters accordingly.

You may also create new parameter groups, and add any combination of measured parameters to this new group. This is explained below.

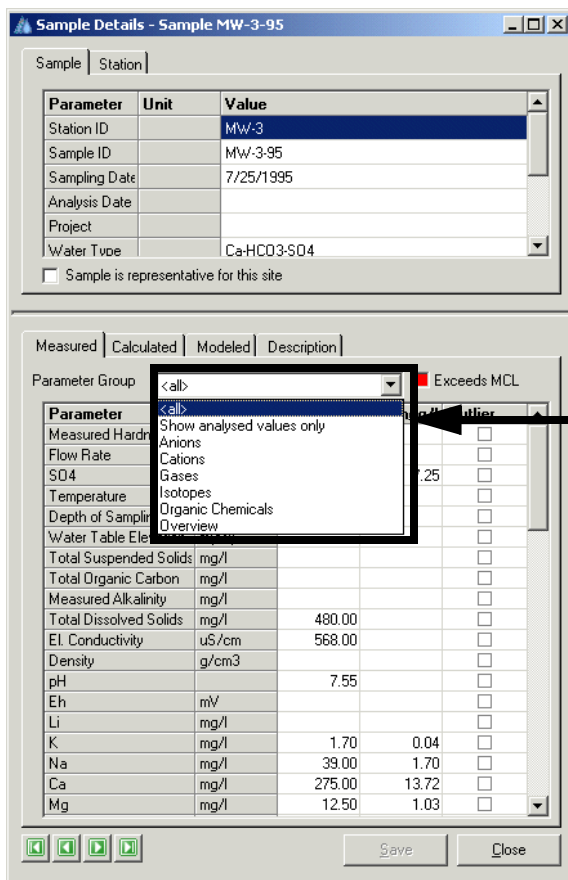
Creating New Parameter Groups

To create a new parameter group:

- Press the  button beside the list of **Parameter Groups**:
- In the **Parameter Groups** dialogue that appears, press the  button to add a new item.
- Double click on this label and type in a name for this new group. (To delete existing parameter groups, select an item then press the  button.)
- Once you are finished press **[OK]**.
- Select this new parameter group from the combo box listing the available groups.
- Use the  button to move parameters from the master list into this group. To move several parameters press the <Ctrl> key while you select multiple parameters from the source list, then press  button. To remove parameters from the parameter group, select the target parameter and press the  button.
- Once you are finished in the parameter groups dialogue, press **[Save]** to save the changes and **[Close]** to return to the main menu of AquaChem.



The parameter groups can now be accessed in the **Sample Details** window. To do so, open the details for any sample and you should see the following window:



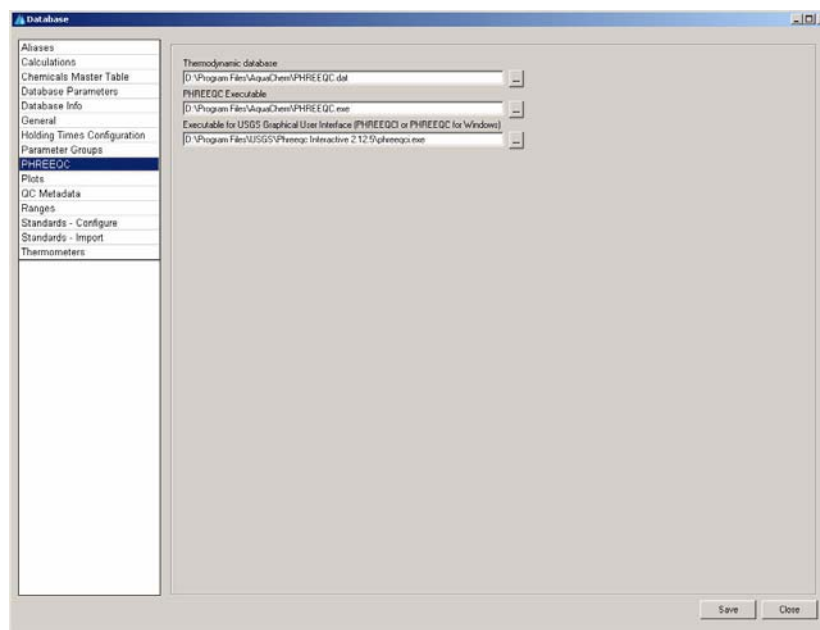
In the **Measured** parameters tab, beside the **Parameter Group** label, there is a combo box listing the available parameter groups for the database (indicated by the arrow). Simply choose one of the groups from this list, and AquaChem will display only the data for the pre-defined parameters in that group.



PHREEQC

This option provides specification of PHREEQC related Files and Folders.

Files and Folders

In this dialogue (shown below), you must specify the locations of the **PHREEQC Thermodynamic Database** (phreeqc.dat) and the **PHREEQC Executable** (phreeqc.exe) files. These files are required for running PHREEQC simulations.

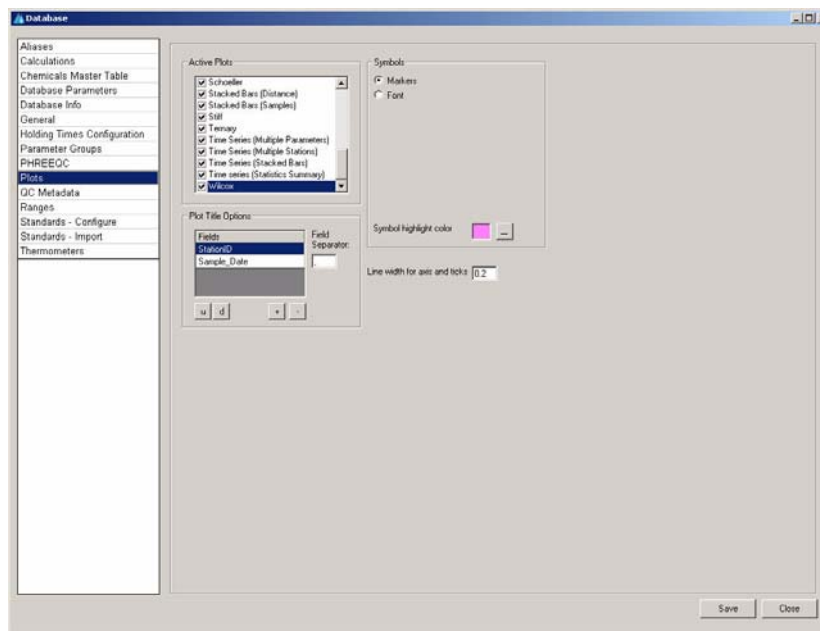


Press  button beside the **Thermodynamic Database** field and select the path and filename from the **PHREEQC Thermodynamic Database** dialogue. From this dialogue, select either **phreeqc.dat**, **Wateq4f.dat**, **Pitzer.dat**, or **Minteq.dat**. In addition, in the **PHREEQC Executable** field select the path and filename for the **phreeqc.exe** file. Finally, specify the path and filename for the USGS graphic user interface (PHREEQCI or PHREEQC for Windows executable, if this component is installed). To change the directories for these files, press the  button beside each directory path, and specify the new file/folder location. By default, all PHREEQC input and output files will be saved in your AquaChem installation folder (default is *Program Files\AquaChem*). PHREEQC-I or PHREEQC for Windows is required for Advanced PHREEQC Simulations.

Once you are finished, click [**Save**] then [**Close**]. You are now ready to run PHREEQC.

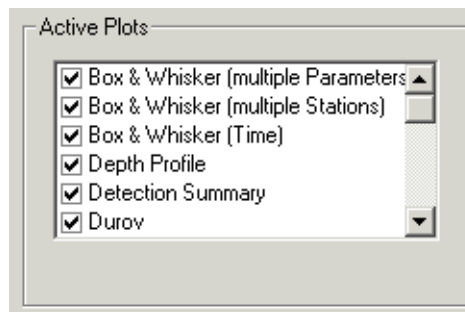
Plots

The Preferences dialogue contains settings for **Plots** which is used for the settings in the Plots and Symbols.



Active Plots


The upper left frame of this dialogue lists the **Active Plots**. This allows you to activate / deactivate the plots. Deactivated plots will not appear in the **Plots** menu (main toolbar). This allows you to hide plots which are never used from the menu. Active Plots are indicated by the presence of a checkmark beside the plot. By default, all plots are active.

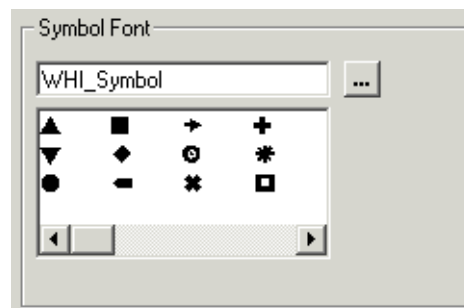


Symbol Font

In the upper right frame of this dialogue you will see options for the **Symbol Font**.

AquaChem provides two options for showing symbols on plots and graphs: **Geometric Markers** and **True type fonts**.

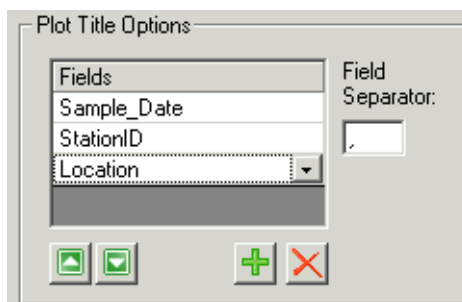
If the Geometric Markers option is selected, you will be able to select from a variety of geometric shapes when defining plot symbols. If the True type fonts option is selected, AquaChem will use “True type fonts” for symbols on plots and graphs. As such, there is an unlimited number of symbols that you can use for your project database. The default font selection is **WHI_Symbol**. To choose another font type, press the  button.





NOTE: The symbols can be defined for samples through the **Plots > Define Symbol or Line** dialogue.

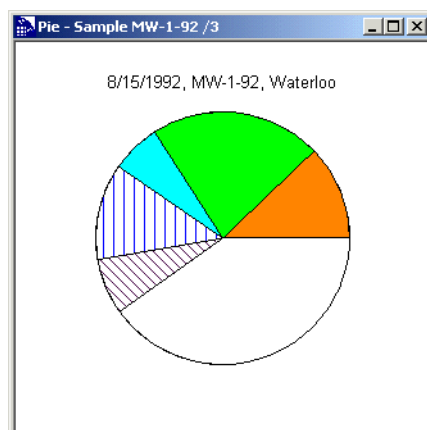
Plot Title Options

This frame includes settings for the automatic plot titling option. When a new **Pie**, **Radial**, or **Stiff** plot is created with your samples, the plots will automatically be titled with the descriptive fields (parameters) listed here. You can specify which parameters you would like to be utilized when using the automatic title option.



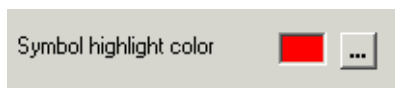
To do so, click the  or  button to add or remove parameters. To change existing fields, simply double click in the field and choose a new parameter from the combo box. The parameter order can be arranged using the up and down buttons. Once you are satisfied with the parameters, you must specify a **Field Separator**. This will be the character that separates the parameters in the title of the plot. You can use any character as the field separator, however the most common are a space, hyphen, comma, or a semi-colon.

Based on these preferences, all new Pie, Radial and Stiff plots will be titled with the unique descriptive parameters for the individual sample. Using these parameters above, you can see an example of a Pie Plot which is created with the automatic title option.



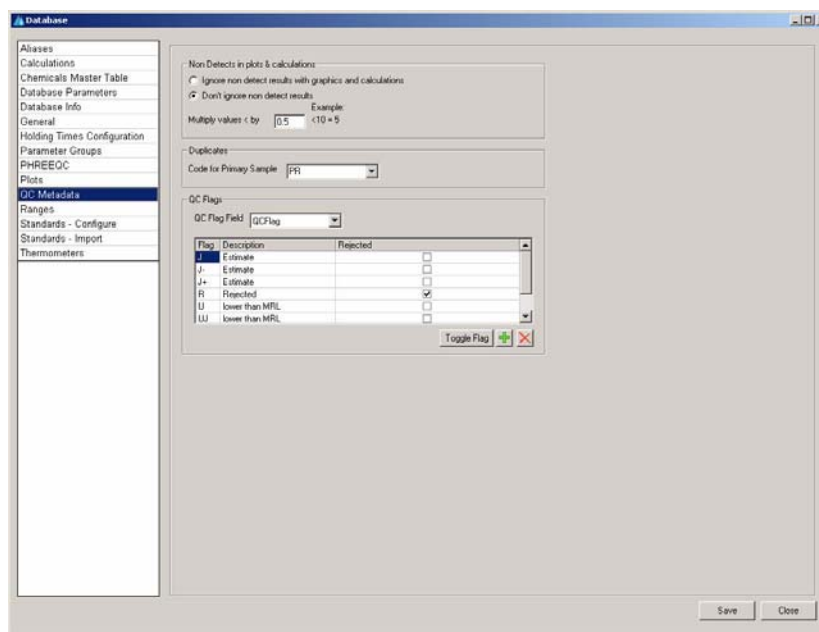
The title of the Pie plot uses Sample_Date (8/15/1992), SampleID (MW-1-92), and Location (Waterloo), separated by a comma.

The **Plots** tab also allows you to choose in which color the selected samples are highlighted when a symbol is clicked. For example, if you use red often as symbol color, you may want to select a different highlight color in order to distinguish which symbols are selected and which ones are not.



QC Metadata

This option allows you to specify settings for several QC analysis options.



Non Detects

Data generated from chemical analysis may fall below the method detection limit (MDL) of the analytical procedure. These measurements are generally described as nondetects, (rather than as zero or not present) and the appropriate limit of detection is usually reported. In cases where measurement data are described as not detected, the concentration of the chemical is unknown although it lies somewhere between zero and detection limit. Data that includes both detected and non-detected results are called censored data in the statistical literature. (Office of Environmental Information, U.S. Environmental Protection Agency, 2000).

In AquaChem database the non detects are entered with a “<” sign. This tab allows you to specify settings for handling censored data in plots and calculation.

In this dialogue, you are presented with options for displaying non detects in plots and all calculations except the Summary Statistics as mentioned above. AquaChem allows you to enter and store approximate measured values using the less than (<) symbols. These options require you to specify whether these approximate values will be ignored or included in the plots.

- If you choose to **ignore non detect with graphics or calculations**, they will be treated as null entries.
- If you choose **Don't ignore non detect results**, you may specify by which factors \diamond these values are multiplied for the use in plots or calculations. For

example, if 2.0 and 0.5 are specified, a value of >10 will be taken as 20, and a value of <10 will be taken as 5.

Duplicates

This option allows you to specify a code to be used to assign to the Primary Sample when a duplicate sample has been found.

QC Flags

This option allows you to specify the Field to be used to populate the flags generated from running a Method Detection Limit Analysis. As well as which flags to be used.

Ranges

Ranges are similar to the drinking water standards; they allow you to define qualitative and quantitative intervals for any parameter. An example for Conductivity (cond) is shown in the dialogue above, and detailed below:


Conductivity (uS/cm)	
100-1000	natural aquifer range
<250	low salinity (C1)
250-750	medium salinity (C2)
750-2250	high salinity (C3)
>2250	very high salinity (C4)

Data can be marked as a “Natural Occurrence” using the provided check-box. If a natural occurrence range is defined for a parameter, the reliability check tool will later be able to detect all concentrations outside this interval. The natural for a parameter occurrence depends strongly of the type of water you are analyzing. It will be very different whether you are working on surface water, ground water or sea water. Please review the values included in the startup database and adjust them to whatever value is applicable in your study area.

Ranges can be useful for interpreting stable isotope data: S, N, C, Sr, He. Isotopes reflect the signature of their source. If the ranges of every potential source is given, an analyzed value can be easily attributed to one of the sources.

Creating a New Range

To add a new Range item:

- Choose the desired parameter from the combo box at the top of this dialogue.
- Press the  button to add a new item.
- Enter the desired **Label** (descriptive name or text).

- Enter the corresponding **Minimum** and **Maximum** values.
- Check the **Natural Occurrence** box if needed.
- Press **[Save]** to save the changes and **[Close]** to return to the main menu.

To view the range options for a sample, load a **Sample Details Window** and then *right-mouse click* on the parameter name. A dialogue titled **Parameter Details** appears, showing all the information about this parameter including the ranges.

Parameter Details - Cond	
Name	Cond
Description	Electrical Conductivity
Unit	uS/cm
MCL	
GCL	
GCL2	
Ranges:	
C1 (Low)	0 - 250
Natural aquifer range	100 - 1000
C2 (Medium)	250 - 750
C3 (High)	750 - 2250
C4 (Very High)	2250 - 10000
Close	

The Range information should appear in the lower section of this dialogue.

Once you are finished in this dialogue, press **[Save]** to save the changes and **[Close]** to return to the main menu.

Standards - Configure

With this option you can specify the set of **Water Quality Standards** to be used for the current database in AquaChem. Multiple standards and levels can be set to active, allowing you to check values against a mix of levels, e.g. ODWS and PWQO, or USEPA and WHO. Values exceeding the respective water quality standards will be highlighted using a color code when displaying the data. You may change the color that should be associated with every level of exceedence by double-clicking on the color for respective level.

Values can also be compared to a percentage of a standard by specifying a percentage (0 -1) under the **Fraction** column for one of the active water quality standards.

The data for each Water Quality Standard is specified in the **Database** dialogue which can be accessed through **File > Database > Water Quality Standards**. There, you may edit existing values for the standards for each parameter. You can also create your own set of Water Quality Standards and import or manually enter values for these new standards.

Viewing Guidelines

Guideline exceedences can be viewed in the sample details table. However, when multiple guidelines are active, there is no legend to find out what exceedences a highlighted cell indicates.

Parameter	Unit	Value	mg/L	Outlier	Comment
pH		7.5			
Potassium	mg/L	2.5	0.06		
Sodium	mg/L	86	3.74		
Calcium	mg/L	125	6.74		
Magnesium	mg/L	23.4	1.93		
Total Iron	mg/L	0.06			
Silver	ug/L	<0.008			
Arsenic (Total)	ug/L	0.5			
Mercury	ug/L				
Barium	mg/L	0.15	0.00		
Lead	ug/L	<0.02			
Chloride	mg/L	128	3.61		
Fluoride	mg/L	0.01	0.00		
Sulfate as SO4	mg/L	205	6.77		
Bicarbonate	mg/L	130	2.13		
Nitrate as NO3	mg/L	<0			
Benzene	ug/L				
Ethylbenzene	ug/L	<2			
Xylene	ug/L	<1			
Toluene	ug/L	10			
Tetrachloroethylene	ug/L	4			
Vinyl chloride	ug/L				
Toluene	ug/L	<1			
Carbon-14	% mod	12.12			
180		-10.15			
Deuterium	I	-73.7			
Tellurium	T.U.	-0.7			

To view all the guideline exceedences for a parameter in the sample details table, simply click on the highlighted cell, and then right-click to display the Parameter Details dialog (shown below).

Parameter Details - Hg	
Name	Hg
Description	Mercury
Formula	Hg
FMW	201.6
Valence	0
Unit	ug/L
CAS Registry Number	7439-97-6
WHO, MCL	1
USEPA Primary DWS	2
Close	


All the active guidelines that are exceeded are displayed at the bottom of the dialog, as indicated in the image above.

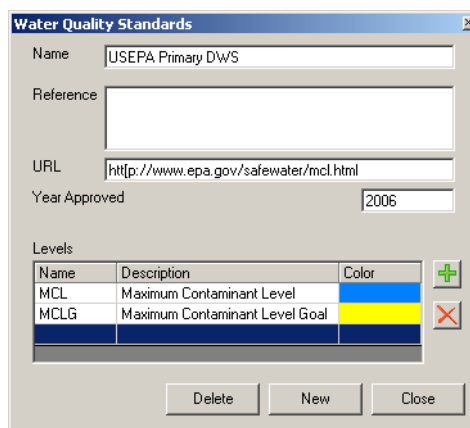
Standards - Import

The options in this dialogue allow you to view and modify water quality standard levels, and create or import new sets of standards. These standards are used in the data analysis reports, table views, and individual sample details. Measured parameters which exceed these pre-defined levels are highlighted in predefined colors depending upon the specified preferences. This allows you to quickly identify sample exceedances and water samples which may be harmful to humans and the environment.

To view the available standards, simply choose a standard from the combo box beside the **Water Quality Standard** field. To view a guideline level for the selected guideline, simply choose a level from the combo box beside the **Level** field.

Standards can be modified by simply editing the existing values beside each parameter in the grid, or importing a complete set of standards from a file (*.txt or *.xls). For each parameter in the database you may define up to three guideline levels.

To view the properties of a Water Quality Standard, press the  button beside the Standard name, at the top of this dialogue. You should then see the following dialogue:




The dialog box titled "Water Quality Standards" contains the following fields and controls:

- Name:** A text field containing "USEPA Primary DWS".
- Reference:** An empty text field.
- URL:** A text field containing "http://www.epa.gov/safewater/mcl.html".
- Year Approved:** A text field containing "2006".
- Levels:** A table with three columns: Name, Description, and Color.

Name	Description	Color
MCL	Maximum Contaminant Level	Blue
MCLG	Maximum Contaminant Level Goal	Yellow

 To the right of the table are a green "+" button and a red "X" button.
- Buttons:** "Delete", "New", and "Close" buttons at the bottom.

In this dialogue, you may view and/or modify the Standard **Name**, **Reference**, and **Year Approved**. It is useful to provide a **URL** to the original standard so that you can check from time to time whether there have been changes made to the standard. The **Levels** frame lists the defined guideline levels and their corresponding colors. You can add a new level by clicking the  button .

You can have an unlimited number of levels for the selected guideline. However, In most cases, two levels will be adequate (Tolerated level, Guideline level). Simply add the number of levels appropriate for the guideline. You may then assign a name to each level (Tolerated level, Guideline level, cleanup level etc.). You can call the first level **MCL** (applicable in U.S.), or you can name this **MAC** (applicable in Canada). Simply type in the new name in the appropriate field. Similarly, for **Level 2** and **Level 3** you may enter your own labels.

For your convenience, several commonly used Water Quality Standards are included with AquaChem:

- World Health Organization (**WHO**) - Guidelines for Drinking Water Quality - 3rd Edition - 2004
- U.S. Environmental Protection Agency (**US EPA**) - National Primary Drinking Water Regulations (EPA 816-F-02-013, July 2002).
- **US EPA** - National Secondary Drinking Water Regulations (EPA 810/K-92-001, July 1992)
- Canadian Council of Ministers of the Environment (**CCME**) - Canadian Environmental Quality Guidelines - 2002 including AO, IMAC, and MAC
- **Health Canada** - Guidelines for Canadian Drinking Water Quality - 2004 including AO, IMAC, and MAC

For more details on these standards, please see the website links below:

WHO: http://www.who.int/water_sanitation_health/dwq/gdwq3/en/

CCME: http://www.ccme.ca/publications/can_guidelines.html

US EPA <http://www.epa.gov/safewater/mcl.html>

Health Canada <http://www.hc-sc.gc.ca/hecs-sesc/water/dwgsup.htm>

If the above mentioned standards are not sufficient for your project needs, you may create a new set of standards, as explained in the following section.

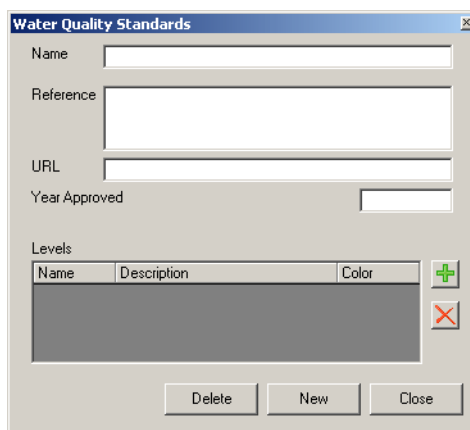
Creating New Water Quality Standards

To create a new water quality standard, follow the steps below:

- If you have not already done so, open the **Database** options dialogue (**File > Database**), and select the **Water Quality Standards** tab.
- Press the **...** button (in the upper right corner) to access the Water Quality Standards properties dialogue, as shown below.

Name	Description	Color
MCL	Maximum Contaminant Level	Blue
MCLG	Maximum Contaminant Level Goal	Yellow

- Click the **[New]** button to create a new standard




A blank **Water Quality Standards** dialog will load (shown above). Here you can:

- Enter the name for this new set of standards
- Enter a reference for the new set of standards
- Enter a URL for the set of standards
- Enter the year that the standards were approved
- Enter guideline levels for the new set of standards (see above for more information)
- Once finished, click the **[Close]** button.

Under the **Water Quality Standards** tab, select the new standards from the **Water Quality Standards** combo box (shown in the image below). Below that, select a guideline level from the **Level** combobox.

There are two ways to enter parameters and assign guideline levels: the values can be entered manually or the data can be imported from a file. Either way, data is entered for one guideline at a time.

Adding Data Manually

To manually add guideline levels, simply click on the  button. A new field will be added to the grey grid, where you can manually enter the following data:

Type	User specified acronym that allows to group parameters, e.g., in chemicals (C), microbial (M) and radionuclide (R) parameters
Parameter*	Parameter name in guideline
CAS#	CAS registry number (for simplified parameter matching)

Unit	Original parameter unit. This may differ from AquaChem's internal unit. The value used in AquaChem may need to be converted.
Standard*	Maximum value or interval for guideline
Comment	A comment
Year	The year when the standard was introduced or revised

**Mandatory Fields*

Importing a Guideline Level


For importing guidelines, an excel file (or text file) must be created for every guideline level, e.g., MCL, MCL goal, etc. The column structure in the excel file must exactly match the column format below, however all columns may be empty **except** the parameter and standard (Upper_Limit) column.

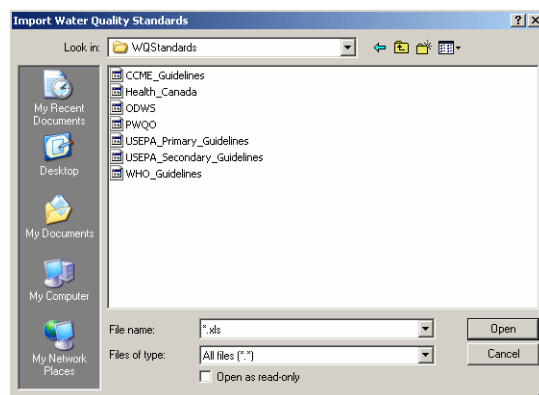
Type	Parameter	CAS	Unit	Lower_Limit	Upper_Limit	Comment	Year
------	-----------	-----	------	-------------	-------------	---------	------

It is recommended that the parameter names in your file are identical to the respective parameter names (the Internal Key name) in the AquaChem database template. If the parameter names are different, AquaChem will not recognize the data for these parameters when you attempt to import them, and you will need to match the fields manually. Optionally, the standard to be imported may contain the CAS registry number per substance, allowing to match this unique code against the CAS registry numbers for the database parameters resulting in a more efficient detection of parameters.

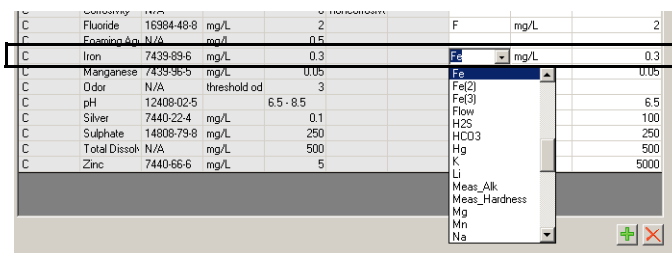
NOTE: Do not include headers in the text file. The first line in your text file should be the first available parameter for which you have a guideline level.

To import the water quality standards using a text or excel file:

- Click the  button beside the **Import water quality standards from File** field for the filename. The following dialogue will then appear:



- Locate the source file on your computer, then click **[Open]**.
- After you load the file, the dialogue will be refreshed
- The grid is divided into two sections: grey fields and white fields. The grey fields represent the original (raw) water quality standard data that was imported. The adjacent white cells represent the matched AquaChem parameters and the corresponding standard levels. The standard values may be adjusted for differences in units between AquaChem and the original source file.
- AquaChem will attempt to identify matching parameters in the database either by using the CAS number if available, or the parameter name. You should carefully scan the list and verify if all the parameters have been identified, e.g. the original name in the guideline may say “Total Dissolved Solids” while the database parameter may say TDS. This will not be discovered by AquaChem and you must select the corresponding AquaChem parameter.
- To manually assign an AquaChem parameter to a standard record, simply click the cell and select a parameter from the combobox (shown below)



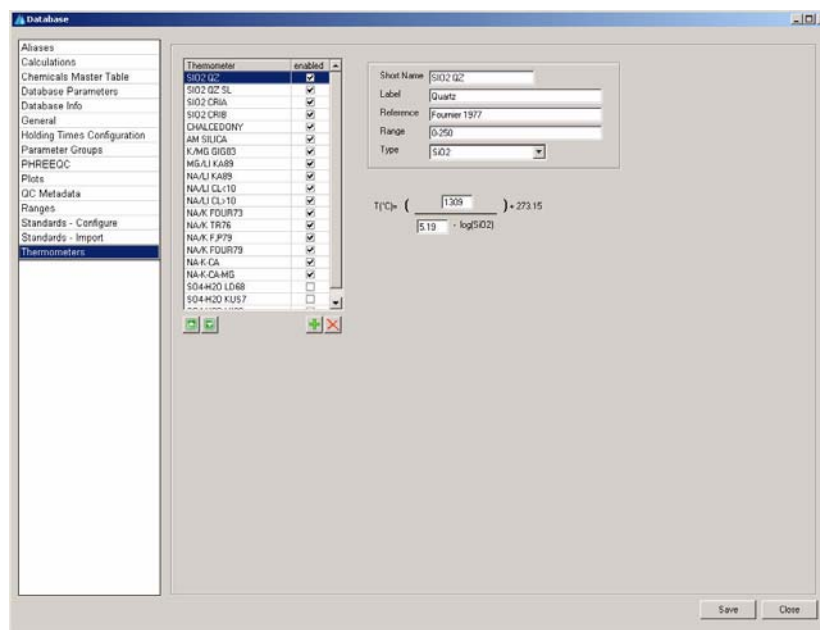
- If the original standard is expressed in mg/l but AquaChem uses ug/l for the same parameter, a conversion factor of 1000 is automatically applied for the value used in AquaChem to check for exceedances. However, for some parameters such as Nitrate and Silica, the parameters may be expressed in different formulas, e.g., nitrate originally as N and AquaChem may express

nitrate as NO₃. In these cases, a conversion factor must be calculated manually and applied to the standard in AquaChem. For example, if the original standard for nitrate as N is 10, then the standard in AquaChem for nitrate as NO₃ is 44.2. You can use the species converter to calculate the conversion factor.

To use this new standard, you must select this in the Standards - Configure option .

Once you are finished in this dialogue, press **[Save]** to save the changes and **[Close]** to return to the main menu.

Thermometers



The Thermometers options allow you to modify existing Geothermometers or to create new ones. Geothermometers are used to estimate the original subsurface reservoir temperature of ascending groundwater which underwent conductive cooling during its ascent. They reflect the temperature dependence of most rock-water reactions. The reactants, which remain dissolved during the journey to the surface, are ideally proportional to their original aquifer temperatures. For example SO₂ (Quartz or Chalcedony dissolution or Na/K for albitisation). The original reference and the temperature range for which the thermometer is valid (most thermometers only work above 100 °C) and can be viewed and edited in this dialogue.

Geothermometers are generally derived from the Van't Hoff equation and have the form:

$$T(^{\circ}C) = \left(\frac{a}{b + \log(K)} \right) + 273$$

where

a and b are constants describing a straight line in the $1000/T\text{-}\log(K)$ plot, and

K depends on the reaction used for the geothermometer (SiO_2 for all SiO_2 thermometers, Na/K for the Na/K thermometer, etc.)

Since thermometers are usually developed from empirical data, different coefficient values are found in the literature. AquaChem includes a compilation of most currently used geothermometers as shown in the figure above. The equations for each of the thermometers are stored in the project database file and can be customized and modified. Each thermometer record includes the formula's coefficients, the reference (author and year), and the range within which the formula was calibrated.

The **Thermometers** tab contains a list of available geothermometers that can be plotted on a geothermometers plot. If a thermometer is enabled (checked) it will be available for the geothermometers plot. If it is disabled (not checked) it will not be available for the geothermometers plot.


There are several fields for each thermometer:

- The **Short Name** refers to the list name of the selected geothermometer.
- The **Label** refers to the reference name of the geothermometer.
- The **Reference** refers to the author and year of the original publication.
- The **Range** refers to the temperature range within which the thermometer is supposed to provide reliable data. This normally corresponds to the range of the calibration data.
- The **Type** refers to the thermometer class.

Creating a New Geothermometer

You may add published thermometers from literature or create your own thermometer if the formula structure corresponds to one of the pre-defined thermometers provided by AquaChem.

For example, to create a new Na/K thermometer:

- Select an existing Na/K geothermometer,
- Click the  button to create a new Na/K geothermometer (new thermometer will be added to the bottom of the list).
- Select the new geothermometer **Type**.
- Enter a new **Short Name**, **Label**, **Reference**, **Range**, and **Type**.

- Replace the formula parameters from the original geothermometer with the parameters from the new geothermometer.
- Press the **[Save]** button to save the changes to the database.

This new thermometer is automatically included in thermometer calculations and can be visualized as a **Geothermometer Plot**.

3.1.12 Exit

The **Exit** command will exit AquaChem. Ensure that all changes are saved to your database before exiting the program.

3.2 Edit Menu

The Edit menu provides access to commonly used Windows tools such as **Cut**, **Copy**, and **Paste**. The Cut, Copy, and Paste commands are not available when the active list window is the active window; these options are only available when you are viewing or editing data for a sample or a station.

3.2.1 Cut

Cuts selected data from the field and places it on the Windows Clipboard.

3.2.2 Copy

Copies the selected data or plot to the Windows Clipboard.

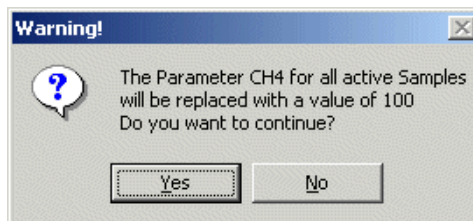
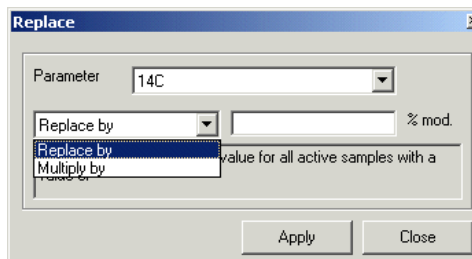
3.2.3 Paste

Pastes data from the Windows clipboard into the selected field. In order to paste numerical values into a sample details field, the field must be active, which is indicated by a flashing cursor in the field. Simply double-click in the appropriate field to activate it and it will be ready to accept data from the clipboard.

3.2.4 Replace

The **Replace** command allows you to replace data for an individual parameter with a new value for all active samples in the database or to multiply all existing values by a specified factor.

Simply choose the method you wish to apply and the parameter you wish to apply this change to from the combo box beside **Parameter**. Then enter the value to replace or multiply by in the field beside the method combo box. Once you are finished, click **[Apply]**. A warning message similar to the one shown below will appear:

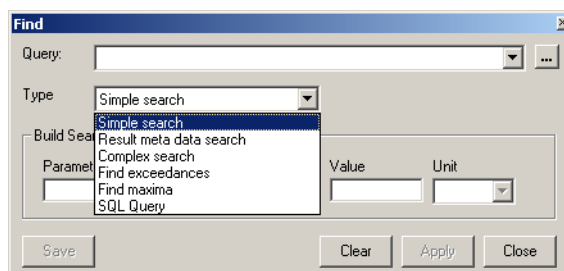


Choose **[Yes]** to proceed or **[No]** to cancel the operation.

The Multiply by method may be useful for correcting import errors. E.g. the X coordinates can be multiplied by 1000 in order to convert the inadvertently imported km values into meters. Note that this method can also be used to bulk edit analysis meta information such as Protocol or Method Detection Limit. For example if you know that in all imported samples, the detection limit of mercury is 1 ng and the method 245.1 was used for analysis, the replace feature is the fastest way for updating this information. Alternatively this information has to be entered manually in the sample detail screen, if it was not included in the import data source.

3.2.5 Find

The **Find** dialogue allows you to create queries of the samples or stations in the AquaChem database.



You may run a **Simple**, **Result meta data**, **Complex Search**, **Find Exceedances**, **Find Maxima**, or **SQL Query** search. Each query type is discussed below.

The records which satisfy the defined search criteria will then be highlighted in your active list. You can now use the filter options in the active list to reduce the list of samples or stations to the highlighted records or to remove highlighted records from this list.

Simple

A simple search allows you to specify one criterion based on Station, Sample, or Measured values.

All records matching the search criteria will be selected (highlighted) in the active list window after the search operation is completed. This feature is very useful for filtering the active list to include only those samples or stations which meet a specific criteria.

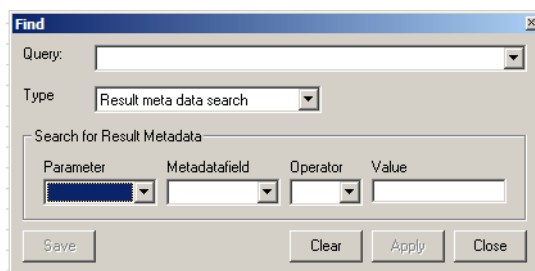
To run a **Simple Search**:

- Enter a parameter in the **Parameter** field, or click **...** button to select a parameter from the list.
- Select the desired **Operator** (<, >, =, LIKE). Searches for strings that use the = operator return exact matches, whereas searches that use the LIKE operator return matches that contain the substring you have specified in the **Value** text field. Searches for character strings are not case sensitive.
- Enter the corresponding parameter **Value**. (The units for the parameter value will be read from the Database options for the selected parameter.)
- Press **[Apply]**.

Result meta data

A **meta data** search allows you to search for result meta data, e.g. all samples where the method detection limit of mercury is above a given value. An example of useful meta information is also whether the results originates from a filtered or unfiltered sample. The meta data search allows you to quickly search for all samples having calcium concentration results from filtered samples.

When you select **Result meta data search**, the following fields will be presented:



The set-up for the meta data search is very similar to that of a simple search, with the exception of the extra field, Metadata field, which allows you to select the appropriate meta data.

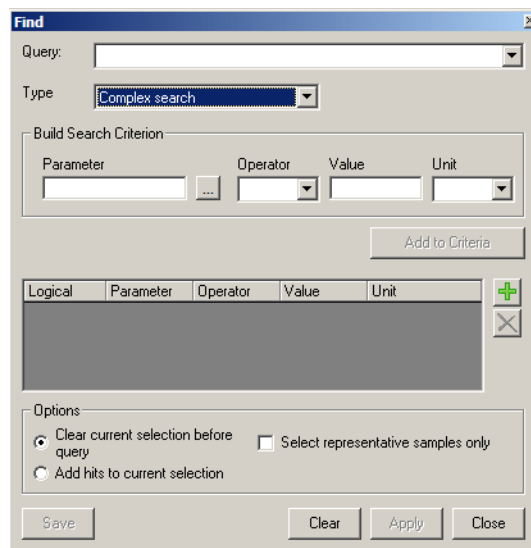
To run the Result meta data search:

- From the **Type** combo box select **Result meta data search**
- Select the desired **Parameter**, **Metadata**, and the **Operator**
- Enter the corresponding **Value**
- Click **[Apply]**

Complex Search

A complex search allows you to specify multiple criteria based on Station, Sample, Measured, Calculated, Thermometer, and Modeled values.

When you select **Complex Search** the following dialogue will appear:





This option allows for a more advanced search of your database. It allows you to combine queries with logical operators, select the unit for each concentration parameter and to search on function results as well. (e.g. Find all samples where Cl <50 AND Hardness > 100).

To run a **Complex Search**:

- From the top of the **Find** dialogue, select the **Complex Search** option
- Enter a search string (**Parameter**, **Operator**, and **Value**)
- Press the **[Add to Criteria]** button
- This will cause the search string to be moved to the lower half of this dialogue. You can now add an additional search string.
- Press **[Apply]** after you entered the desired search strings, to run the query.

The **Logical** field is a combo box containing 'AND' and 'OR'. These logical operators are used to link multiple search criteria. This field is only available after you have entered at least one search criterion. The **Parameter** field is used to specify a database parameter on which to base the query. Below the **Operator** field, there is a combo box listing the available comparison operators (>, >=, <, <=, <>, LIKE, NOT LIKE). In the **Value** field, enter a parameter value for the Find criteria. The **Units** field is used to select a unit system for searches based on chemical parameters or their ratios. This may be useful in the following example: you are searching for samples where concentration of sodium exceeds that of chloride, indicating a sodium source other than halite. The concentrations may be stored in mg/l in the database, but you may specify a search for a Na/Cl ratio > 1 expressed in mmol/l.

The **[Add to Criteria]** button is used to transfer the specified search criteria to the Search Criterion list box. You can also type the query expression directly into the Search Criterion text field if you are familiar with the format. The  and  buttons allow you to add or remove search strings directly and to form the search criteria frame.

Under the **Options**, you must specify the query options for the **Advanced Search**:

- **Clear current selection before query**: this will cause the current search query to be cleared, and run a fresh new query.
- **Add hits to current selection**: the samples that were highlighted from a previous query will remain so.

The **Select Representative Samples Only** option will find only samples that fulfill the current query and are marked as representative in the Sample Details window. This might be useful when comparing samples for different sites and you only want to see a few representative samples per site on plots, reports, etc.

You can assign a representative sample for each site when entering the data in the Sample Details window using **Sample > Representative > On** or checking the **Sample is representative of this site** check box in Sample Details window. If you use the **Select Representative Samples Only** option, only the samples which are representative of each site, and which satisfy the search criterion are returned.

Below are a few examples of the Search Criterion and Search Results:

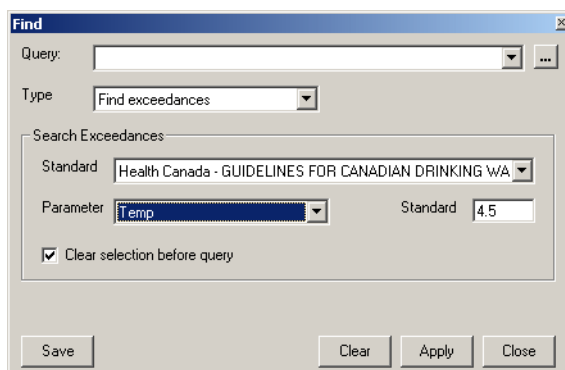
Date > 01/01/1960	Returns all records sampled after January 1st, 1960.
Location = Baden	Returns all records with location "Baden".
Location LIKE Baden	Returns all records with location containing the substring "Baden" (e.g. Baden, Badenweiler, Badentown, etc.)
Na/Cl>1	Returns all records with excess Na with respect to Cl (choose units of mmol/L when using ion ratios).

NOTE: The **Find** selection is context sensitive: if the **Samples** tab is active in the main list, the find will allow you to build and run a Find operation on samples; if the **Stations** tab is active, the find will be applied to stations.

NOTE: AquaChem allows you to save simple or complex queries for easy recall in the future using the **[Save]** button. Saved queries can be retrieved either within the find screen (select from combo box) or in the sample list by selecting it in the **Filter** combo box.

Find Exceedances

A **Find Exceedances** query allows you to find samples with a parameter that exceeds a specified water standard. When you select **Find exceedances** from the **Type** combo box, the following fields will be presented.



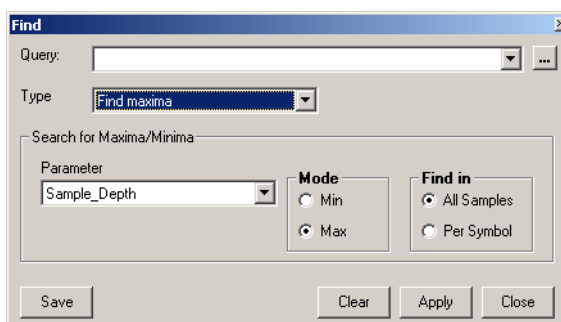
To run a **Find Exceedances** search:

- From the **Type** combo box select **Find Exceedances**
- Select the desired water standard from the **Standard** combo box (See “Water Standards” on page 173 for information on how to add water standards).
- Select the desired parameter from the **Parameter** combo box.
- The corresponding standard value for the selected parameter will display in the Standard text box.
- Click the **[Apply]** button to perform the search.

All records matching the criteria will be selected (highlighted) in the active list window after the search operation is completed.

Find Maxima

The **Find Maxima/Minima** query allows you to find samples with the highest or lowest values of a selected parameter in all samples or just one station.



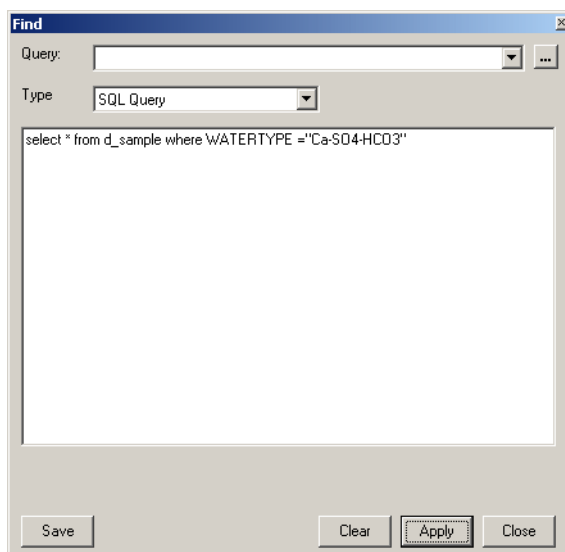
To run a **Find Maxima** search:

- From the Type combo box, select Find Maxima.
- Select the desired parameter from the Parameter combo box.
- Select the desired mode: Min or Max. If you select Min, AquaChem will locate the sample with the lowest value for the selected parameter. If you select Max, AquaChem will locate the sample with the highest value for the selected parameter.
- Choose the group of samples with which to perform the query. Choose **All Samples** to locate the minima/maxima from all samples in the database. **Choose the Per Station** to find to locate the minima/maxima for each station in the database.
- Click the **[Apply]** button

All records matching the criteria will be selected (highlighted) in the active list window after the search operation is completed.

SQL Query

The SQL Query option allows you to query the AquaChem database using SQL syntax. This options requires that you have an understanding of SQL statements as well as good knowledge of the dataset.



To run a SQL Query:

- Type the desired SQL statement in the white space (as shown in the image above).
- Click the **Apply** button.

The query results will be displayed in a separate window, called SQL Query (shown below).

SEL	act	ID	SAMPLEID	STATION_ID	SAMPLE_DATE	ANALYSIS_DATE	PROJECT	SYMBOL	REP	Comment	WATERTYPE	LA
<input type="checkbox"/>	<input checked="" type="checkbox"/>	8	MW-3-92	2	8/8/1992			3	<input type="checkbox"/>		Ca-SO4-HCO3	
<input type="checkbox"/>	<input checked="" type="checkbox"/>	10	MW-3-94	2	6/15/1994			3	<input type="checkbox"/>		Ca-SO4-HCO3	

The query results can be refined further by using the **Find** field in the top left corner of the window. The contents of the window may be copied to the clipboard or sent to a printer by right-clicking on the table, and selecting the desired option.

Find within Distance

The **Find within Distance** command can be found from the main menu under **Edit > Find within Distance**. This command is only available when the **Stations** tab is selected in the active window. Entering a specified distance (metres) in the dialog box of the **Find within Distance** window and clicking **OK** will select all stations in the active window within the specified distance.

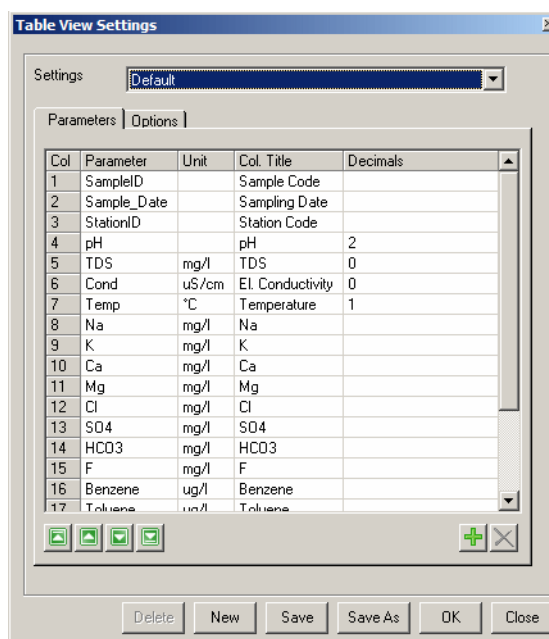
3.3 View Menu

3.3.1 Table View

AquaChem allows you to view and edit data in a tabular spreadsheet-like format containing all samples currently displayed in the Active sample list. This is a convenient method of handling a series of data or comparing selected parameter values for multiple samples. To create a spreadsheet view of the database, select **Table View** from the View menu. You can then use the **Create** option to create your own template, view the **Default** table view, or any customized table view.

Edit

When you select the **Edit** option, a **Table View Settings** dialogue will appear (as shown below) listing the parameters that will be included in the columns of the spreadsheet.



The **Parameters** dialogue tab can be used to select only those parameters you are interested in viewing/editing in a spreadsheet format.



At the top of this dialogue, in the **Settings** field, you will see the name of the current table.

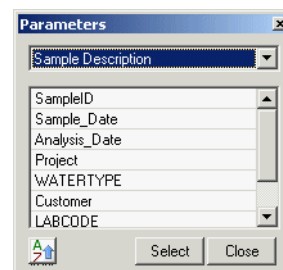
In the lower half of the **Table View settings** dialogue, you will see the list of parameters as they will appear in the table. The order of the parameters can be changed by using the top, up, down, and bottom arrow buttons (as shown here).




Parameters can be added or removed from the table view.

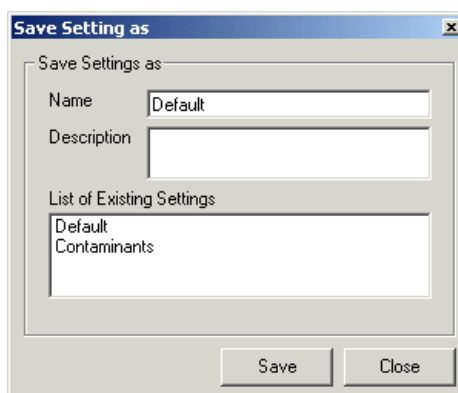
To add a parameter,

- Click the  button in the **Table View settings** dialogue and a **Parameters** dialogue will appear as shown on the right.
- Select the Parameter category by pressing the  button
- Choose parameter(s) from the list that appears
- Press the **[Select]** button.
- Press **[Close]** to return to the previous dialogue.



To delete a parameter(s), simply select parameter(s) from this list and press the  button.

The **Unit** for the table view can also be changed, along with the column (**Col Title**) or row titles and the number of digits used to display the results. Limiting the number of digits is useful when displaying function or ratio results. Once you are satisfied with the Table view, press the **[Save]** button at the bottom of this dialogue. Alternatively, you may save the table view settings under a new name. To do so, press the **[Save As]** button at the bottom of this dialogue, and this will load the **Save Settings** dialogue as shown below.




In this dialogue, enter a **Name** and an optional **Description** for the new Table View you have created and press **[Save]** to return to the **Table View settings** dialogue. Saved settings can now be retrieved from the **Settings** combo box of the **Table View Settings** dialogue. In addition, this new Table View will now appear under the AquaChem **View > Table View** menu below the **Default** table view.

The **Options** tab allows you to further customize the report.

Stat. Param	Title	Value
MIN	Minimum	
MAX	Maximum	
N_EXC	Number of E	

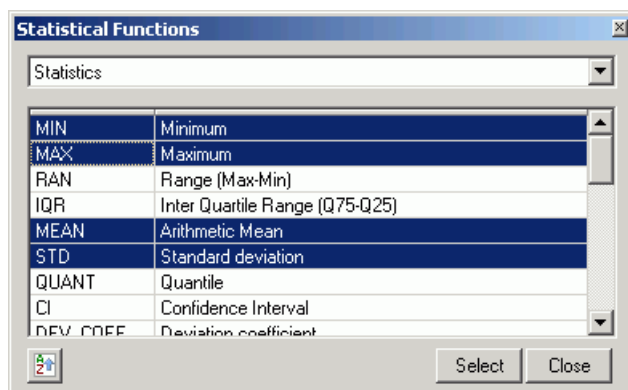
The **Format** may be set to **Samples as Rows** (standard) or **Samples as Columns**; the latter gives a better overview if the spreadsheet contains many parameters, but only a few samples. In the **Highlight Exceedances** combo box you may define whether and how values exceeding the currently active water standard will be highlighted. The selection includes using a bold font for the exceeding values or marking the cell with background color showing which level of the standard has been offended. The legend for those colors is shown at the bottom of the screen. The **Show Standard** option allows you to include the standards values below the row or column of parameter names. Note that all levels of the standards will be shown for multi level standards, e.g. MCL (Maximum Contaminant Level), MCLG (MCL Goal), AO (Aesthetic objective). You can define the code that displays for both non-detect values and not analyzed values, using the **Non Detect Code** and **Not analyzed code** text boxes, respectively. In the **Show Statistics** grid, the user may define statistical values that will be added below the last sample row or column. All summary statistics functions may be used.

Typically the minimum, maximum, mean, percentile, etc., functions are useful in this context. To add statistical functions,

- Press the  button
- Select one or several functions from the statistical functions

- Press the **[Select]** button in the screen shown below

Back in the spreadsheet option tab, you may change the default column or row title for the statistics function using the title column in the screen shown above. For percentiles, you must define the fraction of percentile in the value column that should be calculated, e.g. enter 0.25 for a quartile or 0.5 for the median.



Once you are finished, click **[OK]** to load the **Table View** window.

Spreadsheet [Default]													
Parameter	Mg	Cl	SO4	HCO3	F	Benzene	Toluene	Ethylbenzene	Xylene	Vinyl chloride	PCE	TCE	
Unit	mg/l	mg/l	mg/l	mg/l	mg/l	ug/l	ug/l	ug/l	ug/l	ug/l	ug/l	ug/l	ug/l
CCME MAC						5				2			
USEPA Secondary DWS, MCL		250	250		2								
WHO MCL					1.5	10	700	300	500	0.3	40	70	
CCME AD													
1	22	125	950	125		<1	<1	<2	<1	<1	<2	<2	3
2	25	121	325	150.6	2.3	<1	<1	<2	<1	<1	<2	<2	3
3	23.4	128	325	130	0.01	<1	<1	<2	<1	<1	<2	<2	4
4	26.5	115	275	152.5		<1	<1	<2	<1	<1	<2	<2	10
5	26.2	115	284	144	1	<1	<1	<2	<1	<1	<2	<2	6
6	18.3	100	221.5	145		1	<1	<2	<1	<1	<2	<2	3
7	17.5	111	216.8	126		<1	<1	<2	<1	<1	<2	<2	3
8	19	46	368	463		0	5	<2	<1	<1	<2	<2	<2
9	20.4	42	340	502	2.2	0	2	<2	<1	<1	<2	<2	<2
10	19	65	376	454		0	4	<2	<1	<1	<2	<2	<2
11	12.5	30	348	496	1	0	3	<2	<1	<1	<2	<2	<2
12	11.9	45	340	473		0	5	<2	<1	<1	<2	<2	<2
13	18.2	32	351	580.5		0	5	<2	<1	<1	<2	<2	<2
14	21.6	45	374	601		0	4	<2	<1	<1	<2	<2	<2
15	5	206	150.3	140.3	2.39	50	35	20	4	<1	<2	<2	10
16	23.3	320	319	152.5		50	20	18	5	<1	<2	<2	3
17	20.7	351	308.2	150.1	2.65	10	20	18	4	<1	<2	<2	3
18	21.5	358	302.8	140.8	2.2	3	10	12	5	<1	<2	<2	3
19	24	398	418	135		2	10	10	5	<1	<2	<2	7
20	23	361	299	146		1	10	3	3	<1	<2	<2	3
21	21.4	295	379.2	152.5		1	10	<2	3	<1	<2	<2	25
22	17.6	8.9	732	141.5	<5	<1	<1	5	<1	<1	<2	<2	<1
23	20.6	13	725	144		<1	<1	2	<1	<1	<2	<2	<2
24	20.6	2.8	550	145.2		<1	<1	4	<1	<1	<2	<2	<2
25	22.1	8	590	151.3		<1	<1	3	<1	<1	<2	<2	<2
26	20.2	11	550	145.2		<1	<1	5	<1	<1	<2	<2	<2
27	18.0	3.3	610	126.9	2.03	<1	<1	5	<1	<1	<2	<2	<2
28	19.7	3	604	152.5	2.61	<1	<1	4	<1	<1	<2	<2	<2
29	19.7	3	604	152.5	2.61			10					
30	<1	<1	<1	1.2	<0.2	<1	<1	<1	<1	<1	<2	<2	<1
Minimum	0	0	0	1.2	0	0	0	0	0	0	0	0	0
Maximum	26.9	361	732	601	2.7	50	35	20	5	0	100	10	10
Arithmetic Mean	19.4	116.8	387.8	224.6	1.7	5.9	5.2	4.0	1	0	12.1	3.5	3.5
Number of Exceedences	0	7	26	0	9	6	0	0	0	0	3	0	0

In this dialogue, the colored cells indicate sample exceedences, as defined by the currently selected Water Quality Standards (WHO, CCME or US EPA). Samples which

exceed the guidelines will be shaded with the appropriate color, as defined in the **File>Preferences>Standards - Configure**. At the top of the table view, an water quality standards table is displayed.

You may edit the data in this window as needed; simply edit the required cells with new values. New data will be automatically saved to the database.

NOTE: You may not edit a cell displaying a station parameter. The reason for this is that the station parameters are not entered for each sample; instead each sample has a reference to the station table. Editing a station would edit this station in all samples where it is referenced. If you need to reference a different station for a sample, use the **Assign Station** option when you are in the Sample Details window (**Sample > Assign Station**). If you need to assign a station that does not yet exist in the database, create a new station using the **Station > New** command. You also may not edit values in the cells that are assigned formulas, e.g. a ratio of one parameter to another or the calculated electroneutrality, hardness, sum of cations, etc.

You may print the table view by pressing the **[Print]** button at the bottom of the window, or select **Print** from the main menu.

Select **File > Save** from the main menu to save the table view; the file may be saved in .TXT, .CSV, or .XLS format.

To return to the main AquaChem window, press **[Close]**.


Default

The **Default** command will load a default **Table View** with default parameter settings.

Contaminants



The **Contaminants** command will load a **Table View** with common contaminants.

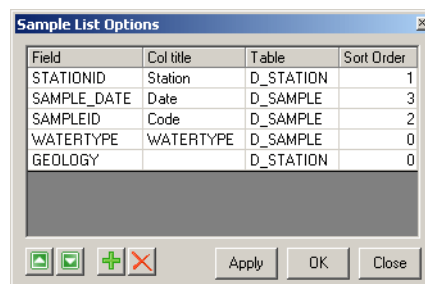
3.3.2 Options

The **Options** command will load the options dialogue for the current AquaChem window. There are options available for the various AquaChem windows. The options will be different for a plot, table view, active list, and the Sample Details window. Each of these is explained below. The same function is performed by the Options  icon on the toolbar.

Options - Active List



The active sample/station list can contain any sample/station description parameter from your database. To edit the parameters in the active list, select **Options** from the **View** menu (alternatively you can press the **[Sort]** button at the bottom of the active list window).

This dialogue allows you to add/remove the parameters which appear in the active list of samples or stations, or adjust their position using the up/down buttons. To add a new item click the  button; to remove an existing item, click the  button. Note that for performance reasons only sample and station parameters can be included in these lists. Additionally you can change the existing parameters. To do so, double-click on a parameter and select a new parameter from the combo box.



The **Col title** column contains the display names for each field. This column is editable, allowing you to modify the display titles of each field.

The **Sort Order** column allows you to specify the sort options for your active list. The active list will be ordered based on the assigned sort number given to each field. For example, in the image above, the list will be sorted by Station_ID and then by Sample_Date and Sample_ID.

To move fields in this dialogue, use the  or  buttons.

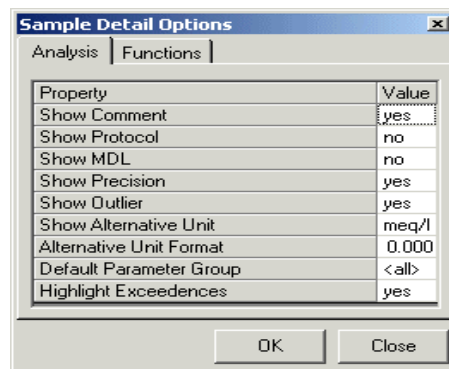
When you are finished press the **[Close]** button at the bottom of the dialogue to apply the changes to the active list.

NOTE: The active samples list and the active stations list each has its own sorting options. To access the sorting options for the Active Stations list, ensure the **Stations** tab is selected in the active list when you select **View > Options**.

Options - Sample Details

When a **Sample Details** window is active, click **Options** from the **View** menu to access the **Sample Details Options** dialogue as shown on the right.

Under the **Analysis** tab, you can edit the settings of the Sample Details window, specifically the **Measured** parameters tab. Fields can be turned on or off, by selecting **Yes/No** from the relevant combo box. You may also select the alternative units, a default parameter group, and enable/disable guideline exceedances (to activate or deactivate this option). You may show or hide the following columns:





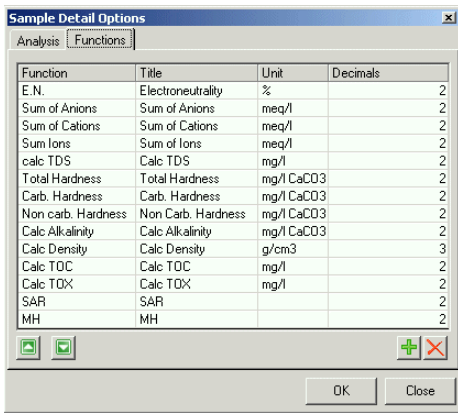
The screenshot shows the 'Sample Detail Options' dialog box with the 'Analysis' tab selected. It contains a table with two columns: 'Property' and 'Value'. The 'Value' column contains dropdown menus. At the bottom are 'OK' and 'Close' buttons.

Property	Value
Show Comment	yes
Show Protocol	no
Show MDL	no
Show Precision	yes
Show Outlier	yes
Show Alternative Unit	meq/l
Alternative Unit Format	0.000
Default Parameter Group	<all>
Highlight Exceedences	yes



- Comment: allows you to add a comment for each measured value (e.g. any difficulties encountered while measuring a particular parameter)
- Standard: select a level within the current water standard to be displayed with every measured result
- Protocol: protocol used for sample measurement
- MDL: method detection limit
- Precision: precision of measurement
- Outlier: outlier check-box (data considered outliers will no be shown on plots or included in calculations)
- Alternative unit: displays an alternate unit (e.g. meq/L when a value is expressed in mg/L)
- Alternative unit format: the conversion process from one unit to another may generate many decimal places. Use this filed to set how many decimal places you want to appear
- Default parameter group: the parameter group that is loaded by default (parameter groups may be defined in **File>Database>Parameter Groups**)
- Highlight exceedences: choose whether or not to highlight guideline exceedences

If one or all of these parameters are not needed, you may hide the respective columns from the Sample Details window.

Under the **Functions** tab (shown on the right) you may edit the settings of the calculations which appear under the **Calculated** tab in the Sample Details window. These options allow you to control the appearance of the Calculated fields. The names of the Calculated parameters can be edited under the **Title** column. The units and number of decimal places for the Calculated values can also be adjusted. The order of the calculated parameters can be arranged using the  and  arrows.



Function	Title	Unit	Decimals
E.N.	Electroneutrality	%	2
Sum of Anions	Sum of Anions	meq/l	2
Sum of Cations	Sum of Cations	meq/l	2
Sum Ions	Sum of Ions	meq/l	2
calc TDS	Calc TDS	mg/l	2
Total Hardness	Total Hardness	mg/l CaCO3	2
Carb. Hardness	Carb. Hardness	mg/l CaCO3	2
Non carb. Hardness	Non Carb. Hardness	mg/l CaCO3	2
Calc Alkalinity	Calc Alkalinity	mg/l CaCO3	2
Calc Density	Calc Density	g/cm3	3
Calc TOC	Calc TOC	mg/l	2
Calc TOX	Calc TOX	mg/l	2
SAR	SAR		2
MH	MH		2

Functions can also be added or removed using the  and  buttons respectively. Once you are finished, press the **[OK]** button to return to the Sample Details window.

For more details on the Calculations performed by AquaChem, please see the **File > Preferences > Calculations** (See “Calculations” on page 90).

Options - Table View

If you select **View > Options** while a **Table View** window is open, this will load the **Table View Settings** dialogue for defining the view of the selected Table (i.e. available parameters, table layout, etc.).

Options - Plots

If you select **View > Options** while a **Plot** window is open, this will load the **Plot Options** dialogue for that plot type.

Options - Reports

If you select **Options** from the **View** menu, while a **Report** window is open, this will load the **Report Options** dialogue for that specific Report type.

3.4 Filter Menu

The options under the **Filter** menu allow you to temporarily “hide” samples, and work with only a few samples or a subset of your database. Omitted samples will be excluded from any plots, tables, reports, or calculations. These options allow you to store several projects in the same database, but focus on specific sites or locations, and produce site-specific plots and reports.

NOTE: The Filter options apply to both the **Stations** and the **Samples**. For example, when a station is temporarily omitted, all samples associated with that station will also be temporarily omitted from the active list.

Show All

Using this command restores all the samples or stations to your active list, e.g. any samples that you omit can be reinstated into the active list by choosing **Show All** (Ctrl-A) option from the **Filter** menu.

Omit All

This option hides all samples or stations that are currently listed.

Show only selected

This option allows you to display only the selected sample(s) or station(s). All non-selected records will be hidden from the active list. When selecting samples/stations in the active list with the mouse, remember that AquaChem supports the standard Windows selection functions: press <Shift> or <Ctrl> to select multiple records.

Omit selected

This option hides the selected sample(s) or station(s) from the active list. These samples or stations are not deleted; they are simply not shown in the active list window, so that you may work with a subset of the database.

Invert Selection

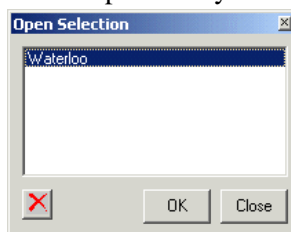
Use this command to select all records that are not selected, and de-select all records that are currently selected.

Select Associated Samples/Stations

Use this command to select all samples which are associated with a station(s) or vice versa. For example, if you are in **Stations** mode and you want to locate all samples which are associated with a selected station, simply click **Filter > Select Associated Samples** then switch to the **Samples** tab and those samples which are associated with that station will be highlighted. This command can be applied to multiple stations or samples. Use the <Shift> or <Ctrl> keys on your keyboard to select multiple records.

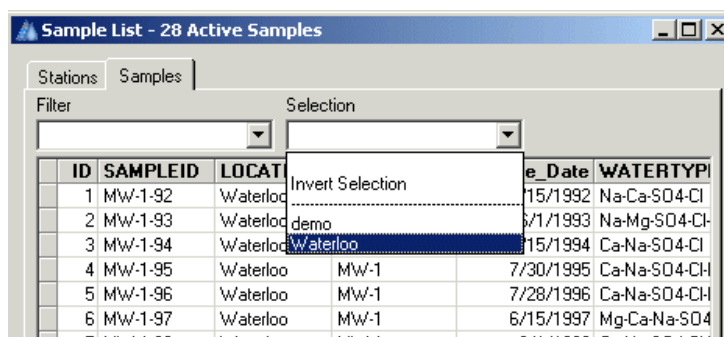
Open Selection

This command allows you to retrieve a previously saved data selection:



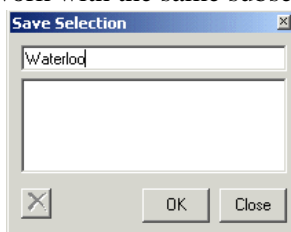
In the **Open Selection** dialogue, select a previously saved selection from the list and press **[OK]**. All records belonging to the selection file will then be highlighted in the active list.

If you want to work exclusively with the samples from the selection file, select **Show Only selected** after opening the selection file. Selections can also be retrieved from the active list as shown below:



Save Selection

The **Save Selection** command saves the list of currently selected samples or stations to your database. In the **Save Selection** dialogue enter the name of the file. This file can be opened in later sessions to work with the same subset of the database.



3.5 Stations/Samples Menu

This menu is automatically updated to reflect the current context (sample or station) of the records list in AquaChem. If the **Samples** tab is selected in your list, or if you have a **Sample Details** window active, then this menu item will show as **Samples**. If the **Stations** tab is selected in your list, or if you have a **Station Details** window active, then this menu item will show as **Stations**. Each of the **Stations/Samples** menu item is explained below.

3.5.1 New

This item allows you to create a new sample or a new station. When you select **Stations > New**, you will see the a **Station Details** window:

Parameter	Unit	Value
Station ID		
Station Name		
Location		
Geology		
Lithology		
X coordinate	m	
Y coordinate	m	
Elevation	m(asl)	
Well Depth		

Navigation icons: [Back] [Forward] [Home] [End] [Print]

Buttons: [Save] [Close]

To add station data, click with your mouse in the input fields and type in the required data. Use the <Enter> key to accept the value and advance to the cell below. Click the **[Save]** button at the bottom of the details window to add the new record to your database.

When you select **Samples > New**, you will see a blank **Sample Details** window:

Sample Details - Sample 0

Sample | Station

Parameter	Value
Station ID	
Sample ID	
Sampling Date	
Analysis Date	
Project	

☐ Sample is representative for this site

Measured | Calculated | Modeled

Parameter Group: Show analysed values only

Parameter	Unit	Value	meq/l
-----------	------	-------	-------

Save Close

NOTE: You must have at least one pre-defined station in your database before creating a new sample. Therefore, when you create a new database, you must first create a minimum of one station; then you may create a new sample and **Assign** this station to this new sample.

To add sample data, click with your mouse in the input fields and type in the required data. Click the **[Save]** button at the bottom of the details window to add the new record to your database.

3.5.2 Clone

Creates a duplicate sample or station. In this procedure only the potentially common data among different samples (such as sampling data, station, etc.) are copied. Cloning samples can be an efficient way of creating multiple samples belonging to the same sampling event.

3.5.3 Edit

Opens the Sample/Station Details window for the selected sample or station.

3.5.4 Delete

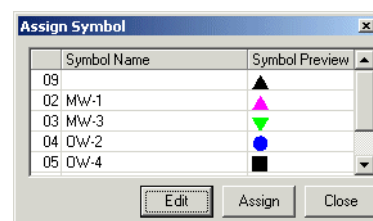
Deletes the selected sample or station.

3.5.5 Assign Symbol

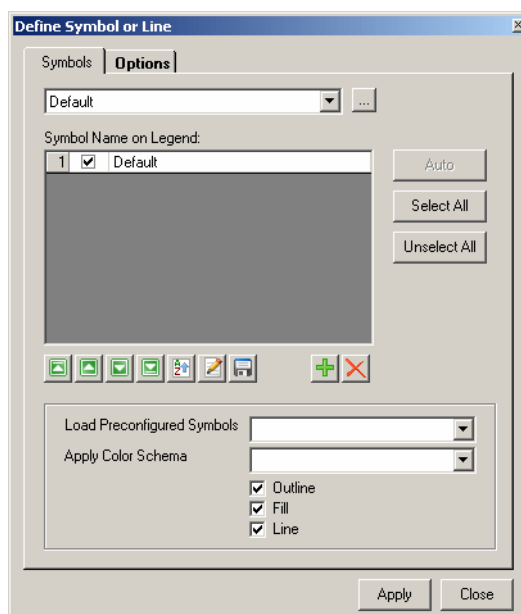
The **Assign Symbol** option allows you to assign symbols to your samples. This option is only available when you are in **Samples** mode.

To assign symbols to samples:

- Select the desired sample(s) in your active list
- From the main menu, select **Samples > Assign Symbol**.
- The **Assign Symbol** dialogue will appear, and provide a list of the available symbol names and the corresponding symbols.
- Choose one symbol from this list and click the **[Assign]** button. This symbol will then be assigned to the selected sample(s).
- To return to the active list of samples press **[Close]**.




To edit the symbol options, press the **[Edit]** button. This will load the **Define Symbol or Line** dialogue as shown below.



In this dialogue, you can create symbols and symbol groups, and define the symbol characteristics. The upper portion of this dialogue contains the **Symbol Group** and the symbols belonging to this group. You can create multiple symbol groups for your samples. For example, you may want to assign symbols based on site, however you may also want to assign symbols based on geology, aquifer body or another variables. To do this, simply create a new symbol group, define the symbols, and assign these symbols to the samples. However, only one symbol group may be plotted at a time. Samples that have not been assigned any symbol of the current symbol group are automatically assigned Symbol 1 of the Default group.



Symbols Tab

There are two default symbol groups included with every new AquaChem database: **Default** and **Station**. The Default symbol group includes one symbol, called Default. The Station symbol group includes one symbol for each unique station ID that is created in your database. Each time a new station is created in your database, a corresponding symbol will be created in the Station symbol group; the symbol name will be identical to the Station ID.

To create new symbol groups, click  button beside the combo box below the **Symbol Group** combo box:



The **Symbol Groups** dialogue will appear as shown below:



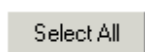
- Click  button to create a new symbol group and then enter a name for the new symbol group
- Click  button to delete existing symbol groups
- Click **[OK]** to return to the previous dialogue

The **Symbol Name on Legend** frame lists the available symbols, their name and status, for the corresponding symbol group.

To add or remove symbols from a symbol group:

- Click  button to add a symbol to the group
- Click on the delete button  to delete the selected symbol

The symbols with a check mark beside them are active; as a result, the samples associated with these symbols will be displayed on the open plots. To remove a group of sample data points from all of the open graphs, simply click on the appropriate check box to de-activate the corresponding symbols. The following buttons also appear in this dialogue:



To activate all symbols, press the **[Select All]** button.



To deactivate all symbols, press the **[Unselect All]** button. To move symbols in this list, use the up/down arrow buttons.



To move the symbol up or down on the list.



To sort the symbols alphabetically.



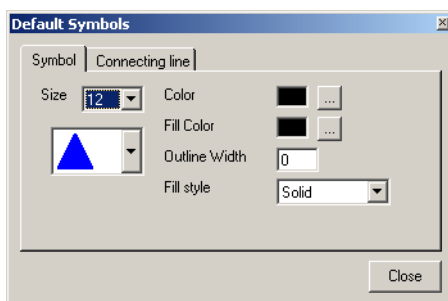
To edit the selected symbol (see “Editing Symbols” on page 158)



To save the current symbol configuration.

Editing Symbols

When the Edit Symbol button is selected, the following dialog will appear on your screen.



Under the **Symbol** tab you will see different shapes of the symbols. The **Color** and **Size** of the symbol can be modified and **Symbol Previewed** as it will appear on the plots. AquaChem is capable of using geometric markers (default) or “True type fonts” for symbols on plots and graphs. The symbol type can be changed in the **Preferences** dialog (see “Preferences” on page 88).

Under the **Connecting Line** tab you can select **Style**, **Width** and **Color**. Each symbol is associated with a particular line style. This line style, the width of the line, and the line color can be customized for each symbol. The **Style** field displays a selection of six line styles, a line width, and a line color that can be displayed in the Depth Profile Plot, Schoeller graph and the Time Series graph to connect associated data points. Line styles other than solid can only be applied if the line width does not exceed 0.3 mm.

Click **[Apply]** to save changes you made to the symbols. To return to the main AquaChem window, press the **[Close]** button.

Options Tab

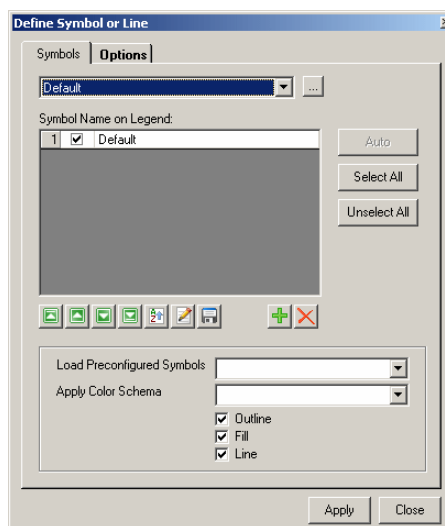
The **Show only symbols present in record list** option allows you to hide symbols, which are not used by any of the active samples. This can be useful when, for example,




your project includes tens of stations but you would like to have quick access to the three stations you are currently working with.

The **Identify Samples** option is used to highlight the samples in the active list that are associated with the selected symbol.

Creating New Symbols

Follow the instructions below to create a new symbol:



- Select a **Symbol Group** from the combo box at the top of this dialogue
- Press the  button and a new field will be added to the list of symbols
- Enter a name for the new symbol
- Then press the **<Enter>** key (on your keyboard) to accept the new name
- Click the **Edit** button to edit the selected symbol
- Choose a symbol character from the list of available symbols
- Choose a **Color** for the new symbol. To access the color options, press the  button beside the color field, and select a color from the Color palette.
- Choose a **Size** for the symbol. A list of font sizes can be accessed by clicking the  button from the combo box.
- Click the **[Close]** button
- Press **[Apply]** to create the symbol
- Press **[Close]** and return to your sample list

Now use the **Samples > Assign Symbol** option to assign this symbol to the desired samples. If done successfully, the active samples window will display a column listing the symbol number which is assigned to each sample.

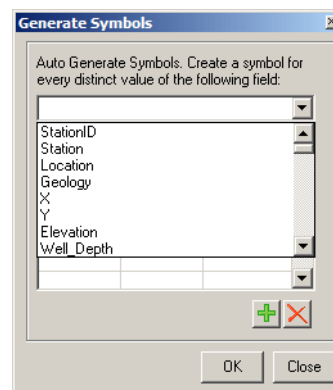
Auto Generate Symbols

AquaChem includes a convenient feature that allows you to automatically create symbols based on station database parameters. For example, your station data contains a parameter “Geology” and ten distinct geology terms have been assigned to the stations within the current project. Instead of searching for all different geology terms and adding a symbol for each one, you may use the **Auto** feature to create these items. Press the [Auto] button in the **Define Symbol or Line** dialogue to load this option. Typically, parameters that should later be used as symbol labels should be defined as list of values (pick list) in order to guarantee consistent spelling.

NOTE: The auto generate symbols option is not available for the Default symbol group. You must first create a new symbol group in order to activate the **[Auto]** button and to use this option.

In the **Auto Generate Symbols** dialogue, you will be prompted to choose a parameter for which symbols will be automatically generated. The available parameters are all Sample and Station description parameters and measured parameters as defined in your database template.

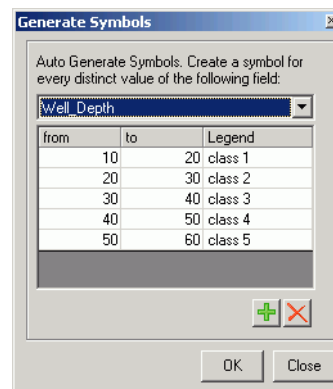
Numeric and text parameters will be handled differently. If you choose a parameter of type text, e.g. Geology, the grid below the selection box will be filled with all geology expression that could be retrieved from the current data set. You may edit or remove expressions that are not required or add new geology expressions. Note that for new expressions, a symbol will be created, but no samples will be initially assigned to this symbol.



After selecting a numeric parameter (any station, sample or measured parameter) the grid below the selection box will be filled with 5 classes dividing the range in which the selected parameter is represented in the database into regular intervals. You may remove, edit and add any intervals and edit the interval legend.

A unique symbol character will be generated for each symbol group, and will use the default font size of 12 and default color black.

Select the desired field and click **[OK]** to return to the **Define Symbol or Line** dialogue, where you will see



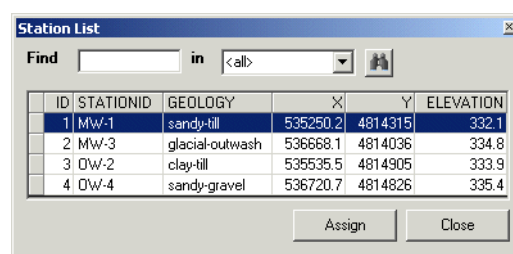
the new symbols which were automatically generated. You now have the option of modifying these new symbols (i.e. the shape, color, name, size, etc.).

The new symbols are automatically assigned to the samples in your database; the symbols will be assigned to the matching parameter for that sample. For example, if you automatically generate symbol groups based on Location, AquaChem will search your database for all instances of the Location parameter, and create a symbol for each unique entry for Location. For numeric parameter, each sample will be assigned to the symbol, where the sample parameter lies between minimum and maximum value specified for this symbol.

3.5.6 Assign Station

The **Assign Station** option allows you to assign a station to the selected sample. This menu item is only available when a **Sample Details** window is active.

When you select **Samples > Assign Station** from the main menu, a **Station List** dialogue (as shown here) will appear with a list of the available stations in your database.



From this dialogue, you may select a station directly from the list; or if you have large list of stations, the **Find** feature at the top of this window can be helpful. Once you have located the desired station for this sample, press the **[Assign]** button at the bottom of this dialogue and this will return you to the Sample Details window.

3.5.7 Representative On / Off

This option defines the **Representative** samples flag for the selected sample(s). A representative sample is indicated by the presence of a check mark in the **Active Samples** window as shown below.

ID	Station	Date	SampleID	WATERTY	GEOLOGY	SYMBOL	REP
1	MW-1	8/15/1992	MW-1-92	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
2	MW-1	6/1/1993	MW-1-93	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
3	MW-1	6/15/1994	MW-1-94	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
4	MW-1	7/30/1995	MW-1-95	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
5	MW-1	7/28/1996	MW-1-96	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
6	MW-1	6/15/1997	MW-1-97	Ca-Na-SO4	silty-gravel	1	<input checked="" type="checkbox"/>
7	MW-1	8/1/1998	MW-1-98	Na-SO4-CH	silty-gravel	1	<input checked="" type="checkbox"/>
15	MW-2	8/1/1992	OW-2-92	Na-Ca-Cl-SC	silty-sand	1	<input checked="" type="checkbox"/>
16	MW-2	6/5/1993	OW-2-93	Na-Ca-Cl-SC	silty-sand	1	<input type="checkbox"/>
17	MW-2	6/12/1994	OW-2-94	Na-Ca-Cl-SC	silty-sand	1	<input checked="" type="checkbox"/>
18	MW-2	7/21/1995	OW-2-95	Na-Ca-Cl-SC	silty-sand	1	<input type="checkbox"/>
19	MW-2	7/24/1996	OW-2-96	Na-Ca-Cl-SC	silty-sand	1	<input type="checkbox"/>

This feature is convenient when handling a large dataset consisting of many samples from multiple stations. By marking manually one or several samples per station as representative, you may later filter these samples much more quickly when creating plots comparing all stations, where the selection of all samples would make the plot confusing.

When you perform a search or calculation, you may choose to use only samples in the active list which are representative of each site.

- To enable this option, choose **Sample > Representative > On**.
- To disable this option, select **Sample > Representative > Off** (or manually remove the check mark in the Sample Details window).

NOTE: The Representative menu item is only available when you are in the samples mode.

3.5.8 Goto next Selected (CTRL+S)

Use this menu item to navigate highlighted records in the sample or station tables.

3.6 Plots Menu

The following section provides a brief summary of the commands in the **Plots** menu. For more details on the AquaChem plots and the plot options, please refer to Chapter 4.

3.6.1 New

This item allows you to create a new plot. See Chapter 4: Plots for more details.

3.6.2 Open Configuration

Opens a previously saved plot configuration.

3.6.3 Save Configuration

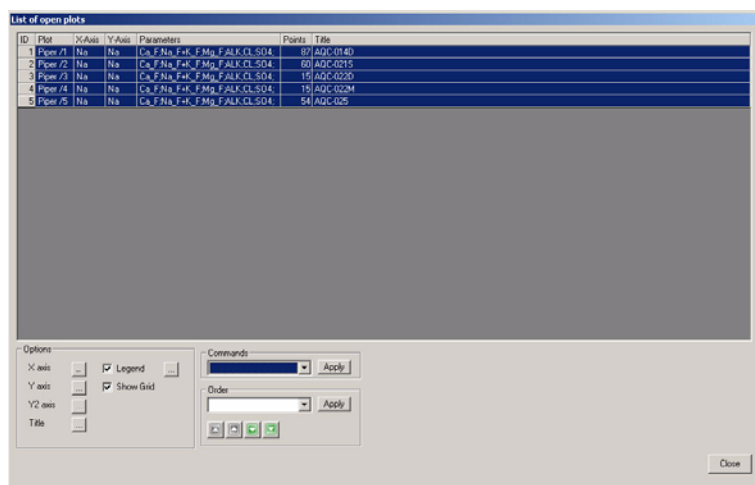
Saves the current set of plots and their settings to the current AquaChem database.

3.6.4 Close All Plots

Closes all plot windows.

3.6.5 List of Plots

Loads the **List of Plots** window. The **List of Plots** contains a list of all opened plots , and allows you to make changes to multiple plots simultaneously. For example, you may have generated time series plots for each station in your project. Using the list of plots, you can quickly adjust the axis extents, change fonts and symbols, etc. for all opened plots, simultaneously.



To make changes to multiple plots in the plots list,

- Highlight the desired plots by holding the **SHIFT** key on your keyboard, and selecting each plot.
- Select one of the buttons in the **Options** frame to load the associated options. You may change settings for the X-axis, Y-axis, Y2-axis (if applicable) plot Title and plot Legend. You may also show/hide the Plot grid. For more information on these plots options, please refer to “Common Plot Features” on page 178.

- Once the desired changes have been made, click the **Refresh** button to apply the changes to the highlighted plots.

You may also select from the following commands:

- **Close All Plots** - closes the highlighted plots
- **Minimize Plots** - minimize the highlighted plots
- **Restore Plots** - displays any of the highlighted plots that have been minimized.
- **Refresh All Plots** - applies changes made in the options to all highlighted plots.
- **Clone Plot** - to make a duplicate copy of the highlighted plot.

You may also select from the following order by options:

- **Plot Title then Y axis Title**
- **Y axis Title then Plot Title**

You may also use the green up and down arrows to order the plots. This information is saved within your plot configuration so when recreating plots you can save time.

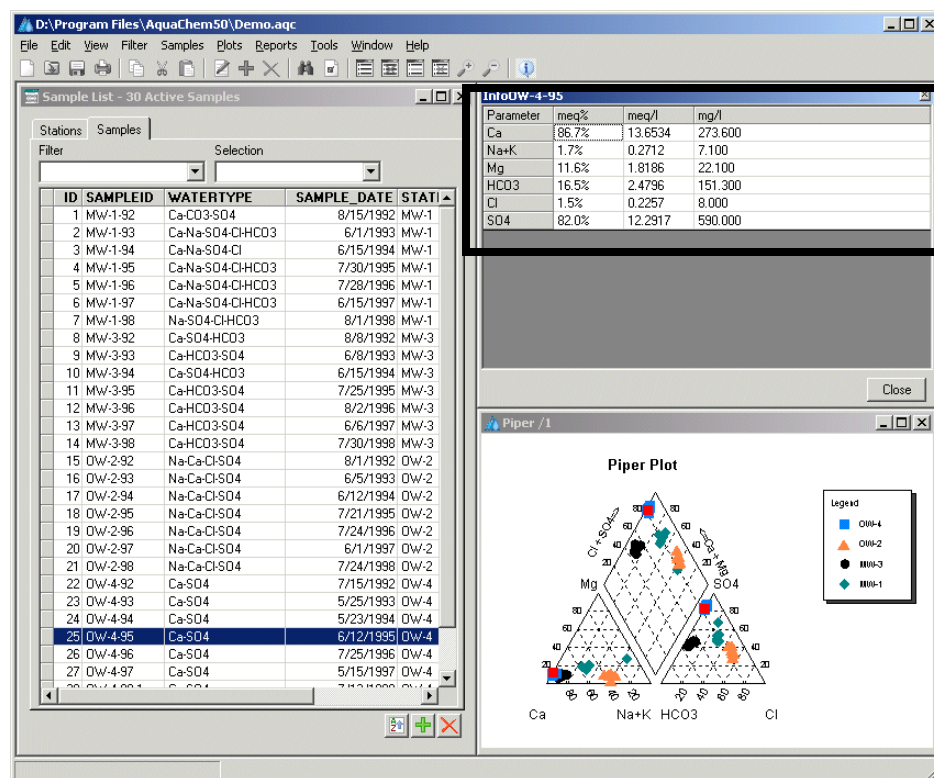
3.6.6 Define Symbol or Line

Loads the **Define Symbol or Line** dialogue for defining symbol and line properties as they appear in the plots and graphs. For more details on this dialogue, please refer to the sections Assign Symbol, Creating New Symbols, and Auto Generate Symbols above.

3.6.7 Show Sample Data

If this option is selected, AquaChem will display a window with the data for the selected sample. Simply click on a sample point from the plot and the **Info** data dialogue will appear. Each time you click on a new sample point, the new data is loaded into dialogue. An example is shown below.

NOTE: You may need to re-arrange the positions of your windows in order to see the sample data dialogue.



3.6.8 Identify Plot Data

The Identify option allows you to link the plotted data to the samples in the active list. This allows you to click on a point in the graph and view the corresponding sample in the active list. This is useful when you have a large number of samples plotted on one plot, and you want to identify outliers.

There are three options available for selecting and identifying points on a plot: **None**, **Selected Plot**, and **All Plots**. The default mode to identify points is on All Plots; this means that all data points on all plots are linked to the database. These three options are described below:

None

If you click on a plot, no samples will be highlighted. The data points on the graphs are not linked to the database which means that samples are not identified. When you have several open graphs and a large database with many samples, it may be convenient to have no link between the graphs and the database in order to increase the speed of the various AquaChem operations.

Selected Plot

Only the data points in the active graph are linked to the database. The plotted point is activated (turns red color by default) and is selected in the sample list. This mode is useful if you are working with a large data set and want to focus on just one graph.

All plots

The sampling point of the corresponding sample is activated on all plot windows and on the sample list. If you are working with several graphs, you can simultaneously identify a sample in all plots by clicking near a point in any of the plots, or by clicking on a sample in the active list. The sample will become selected in the active list, and the corresponding data points will be highlighted in red (by default) in all the open plot windows. If the Table view is open, the records will be identified here as well.

3.7 Reports Menu

The **Reports** allow you to create a summary of your data from your AquaChem database. AquaChem generates seven pre-defined reports and also allows you to create your own report templates.

These reports are all generated in a separate Report window in a spreadsheet view. The reports can be printed ‘as is’ using the **[Print]** button on the lower-left corner of the window or saved using the **[Save]** button.

The following is a brief summary of the options under the **Reports** menu. For more details on Reports, please refer to Chapter 5: Reports.

3.7.1 Data Summary

The **Data Summary** report allows you to generate a summary report for the AquaChem database and all stations and samples currently displayed in the Active Stations/Samples list.

3.7.2 Compare Sample

The **Compare Sample** report allows you to compare a sample in your database to another sample or multiple samples. The Compare Sample report uses a linear regression algorithm to generate the **Correlation coefficient** and the **Euclidean distance** between a selected sample and all other active samples. Samples having a chemical composition similar to the selected sample will have a correlation close to 1 and a small euclidean distance. Samples having similar chemistry, but being diluted with respect to each other will have correlation close to 1, but a large euclidean distance. Samples with little similarity will have a correlation coefficient close to 0.

3.7.3 Mix Samples

The **Mix Samples** report generates solution concentrations resulting from the step-wise mixing of specified proportions of two selected samples from the project database.

3.7.4 Water Quality Standards

The **Water Quality Standards** report provides a summary of parameters exceeding the established Water Quality Standards for the selected sample. The selected samples are compared to the pre-defined Water Quality Standards as specified in the **Preferences** in the **File** menu.

3.7.5 Hardness Dependent Standards

The United States Environmental Protection Agency (EPA) has established standards for metals where toxicity is a function of hardness. The water quality criteria for metals can be expressed as “total recoverable” or “dissolved” for acute and chronic concentrations. The total recoverable also includes the sorbed concentration on particles. Acute standards offer protection from toxic effects from chemical concentrations during a shorter term exposure. Chronic standards offer protection from toxic effects from a chemical during long-term exposure.

EPA Freshwater standards are mainly used for hardness dependent standards, however similar standards can be found in different countries. The demo_basic database provided with AquaChem only includes the EPA Freshwater Standards, however you may create your own standards.

3.7.6 Rock Source Deduction

This report allows you to determine with what kind of minerals the water sample has been in contact with during its subsurface journey. The results are based on ion ratios and total dissolved solids found in the sample. If results do not meet expectations, they should be confirmed with more a detailed study based on multiple samples, aquifer mineralogy analysis, modeling, and plots.

3.7.7 Statistics

The **Statistics** submenu provides several statistical analyses you may wish to perform on your data.

Summary Statistics

Generates the summary statistics such as Minimum, Maximum, Mean, and Percentiles for all samples available in the project.

Correlation Matrix

Generates a correlation matrix for a specified number of sample parameters that are common to all active samples. A linear regression routine calculates the regression coefficient (r), and the slope and intercept of the regression line.

Trend Analysis

Calculates the trend for a selected parameter using linear regression, Sen's test, or Mann-Kendall test.

Outlier Tests

Uses Dixon's (extreme value) test, Discordance test, Rosner's test, and Walsh's test to determine whether there are any outliers for a specified parameter in the dataset.

Tests for normality

Use the Studentized range test (t-test), Geary's test, Shapiro Wilk test (W test), or Skewness and Kurtosis test to determine if the values for a specified parameter follow normal distribution.

Alert Levels

An **Alert Level (AL)** refers to an upper tolerance level for a constituent which can be used as an early warning indicator of a potential violation of a water quality standard. In AquaChem, Alert Levels can be calculated for one or more constituents using baseline monitoring data. The alert levels may then be compared with future detection monitoring data to determine statistically significant increases of constituents in groundwater.

3.7.8 Sample Summary

The **Sample Summary** report provides a general overview of a sample including major ions, hydrochemical facies (e.g. Na-Cl), calculated hardness, ion balance, ion ratios, etc.

3.7.9 Report Designer

The **Report Designer** allows you to customize your own reports based on a single sample. The user defined reports may contain any parameter, ratio of parameters or function values. It is also possible to add a template and create a print ready output for your report. For more information refer to Chapter 5: Reports.

3.8 Tools Menu

The **Tools** menu provides access to the data analysis tools, including converters, calculators, LookUp Tables, and access to the PHREEQC modeling interface. The following is a brief summary of the options under the **Tools** menu. For more details on Tools, please refer to Chapter 6: Tools.

3.8.1 Calculators

When you access the **Calculator** option from the Tools menu, you will find the following options:

Aggregate Samples

This tool allows you to create a new sample based on a number of selected samples, which are aggregated using various algorithms such as minimum, maximum, median, geometric mean, etc.

Aquachem Function

The **Aquachem Function** is a complement to the Calculated parameters tab available in the Sample Details window. It allows you to calculate any function in any available unit quickly and easily, based on the selected sample. The calculations in the Sample Details window are fixed and typically contain only the most commonly used calculations. However, if you want to use some of the other calculations the **Aquachem Function** dialogue provides these options.

Calculate facies

This command calculates the **Watertype** expression for all highlighted samples.

Corrosion and Scaling

This tool estimates a samples tendency for scaling or corrosion

Decay Calculator

This tool allows you to calculate the degradation of a species by a first order reaction:

$$\frac{dC_A}{dt} = -C_A$$

You may choose to calculate:

- **Concentration after a specified amount of time**
- **Time to reach a specified concentration**
- **Estimate half life from given measurement concentration values**

AquaChem includes half-lives for more than 300 species, for physical phases including air, soil, groundwater and surface water. These half-lives are taken from the following reference: Ph. Howard et al. (1991): Handbook of Environmental Degradation Rates.

Find Missing Major Ion

This tool helps you to fill in data for missing cations or anions, using the charge balance and existing measured values for other ions for the sample.

Formula Weight Calculator

This tool helps you to calculate the formula weight based on a user-defined formula.

Oxygen solubility

This command allows to calculate the theoretic solubility of oxygen based on the temperature and elevations of using the American Society of Civil Engineers' formula.

Volume Concentration Converter

This converter is used for samples with measured organic chemicals. It allows you to convert measured concentrations in ppm by volume (ppmV), to mg/m³ at a specified temperature.

Retardation

This tool allows you to calculate the retardation coefficient by entering your porosity, organic content and the K_d (distribution coefficient).

Special Conversions

This tool provides conversions for non-linear geochemical calculations. E.g. Conductivity \leftrightarrow Resistivity, pe \leftrightarrow Eh, etc.

Species Converter

This tool allows you to convert any species into a different form. This is useful to express a measured amount of a parameter as a different aqueous species in mg/L. For example, you may convert Nitrogen as NO₃ into Nitrogen as N or vice versa.

Unit Calculator

This tool performs basic unit conversions for length, time, volume, density, mass, etc.

UTM Conversion

This function allows you to calculate geographic coordinates from projected (UTM) coordinates, and vice versa, for stations in your database.

3.8.2 QA/QC

Compare Duplicates

This report allows you to compare any two records with matching DuplicateID's and determine where they differ for each concentration result.

Control Intervals

This tool allows you to configure your control intervals which can be used when determining if a sample is an outlier.

Highlight Duplicates

Select this option in order to highlight all samples marked as duplicates. Aquachem considers a sample as duplicate, if the Duplicate_ID value of the sample description is not empty.

Highlight Outliers

Selecting this option will load a dialogue in which you can choose a parameter. Clicking **OK** in that dialogue will highlight all samples in the Active Samples list that have the outlier flag selected for the parameter value.

Holding Times

This tool allows you to calculate Holding Times for selected parameters. You must specify which fields contain the date and time used for calculating the holding time under File>Preferences>Holding Times Configuration.

Manage Duplicates

This tool allows you to find and mark duplicate samples by assigning duplicate codes.

Method Detection Limit

This tool allows you to check your samples against Method Detection Limits which are configured by selecting File>Preferences>Database Parameters.

Reliability Check

The **Reliability Check** report allows you to confirm the validity of the measured sample data. There are a number of tests provided in AquaChem which can provide insight into the reliability of your analysis.

Total/Dissolved Comparison

This tool allows you to compare total (or unfiltered) and dissolved (or filtered) concentrations.

3.8.3 Look Up Tables

The **Lookup Tables** shows the tables that are defined in the **File > Database > LookUp Tables** dialogue. These tables allow you to quickly find information for use in AquaChem.

Degradation Rates

This look up table provides a list of degradation rates for common organic chemicals.

PHREEQC Phases

This look up table provides a list of commonly used minerals including formula and formula weights from the PHREEQC thermodynamic database. If no PHREEQC options are specified in the preferences, then this table will be empty.

Periodic Table

This table provides information for the elements of the periodic table.

Water Standards

This table contains all of the water quality standards and guidelines for the current database.

Time Series

This allows you to select and view any table of time series data that you have imported.

Browse Database

This allows you to browse through the entire AquaChem database and view any table.

3.8.4 Modeling

AquaChem has a built-in link to the PHREEQC geochemical modeling program, that is capable of creating one or more solutions from the water quality samples in your AquaChem database. The **Modeling** option under the **Tools** menu provides links to the various options for PHREEQC. The following is a brief introduction to the modeling features included with AquaChem. For more details, please see Chapter 6 and 7.

Calculate Saturation Indices and Activities

This command will run PHREEQC for the samples selected in your active list. PHREEQC will calculate saturation indices and activities for those parameters which are defined in your database (those parameters listed in the Modeled Parameters tab). The results will automatically be saved back to the Sample Details table, for the selected samples.

Calculate pH

This allows you to calculate the pH for the selected samples. The simulation is based on the assumption that the solution is in equilibrium with a carbonate mineral. This utility may be used to estimate the pH, in the case where a measured pH value is not available. However, this is calculated based on the assumption that the groundwater sample in question flows in a carbonate aquifer and is in equilibrium in respect to calcite or another specified carbonate material.

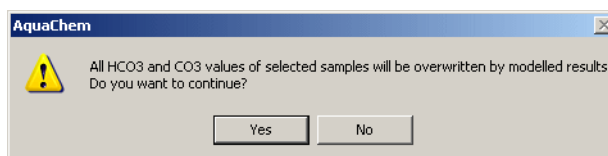
Calculate Eh

This allows you to calculate the Eh for the selected sample based on a redox couple. The Eh is calculated based on the redox speciation, using the Nernst Equation. Each Redox couple gives an individual Eh value which in cases of equilibrium conditions should be reasonable. This menu item launches PHREEQC to search for available redox couples and to calculate pH and Eh values for each couple.

Alk > HCO₃, CO₃

Alkalinity is the measure of the total acid-neutralizing capacity of the water. Since most of the alkalinity in a natural groundwater is due to inorganic carbon species, it can be used to determine bicarbonate and carbonate concentrations. This utility allows to calculate the bicarbonate and carbonate concentration based on the total alkalinity and pH of a selected sample. For a given alkalinity, the bicarbonate/carbonate ratio increases with higher pH values and decreases under low pH conditions.

When this option is selected, the following warning message will appear:



Click [**Yes**] to calculate the HCO₃ and CO₃ modelled values for the selected sample. Any existing HCO₃ and CO₃ values in the selected sample will be overwritten with the modeled results.

Equilibrate with Minerals

This calculator allows to equilibrate a solution with one or several minerals and calculates the amount of mineral that is necessary to reach equilibrium. This equilibration can be combined with a partial evaporation of the sample to simulation the evaporation process. As a further option, a pump rate and duration can be specified and the program then calculates the total amount of solids, that will be precipitated or dissolved during this period.

PHREEQC (Basic)

This allows you to create an input file for PHREEQC, and loads a graphic user interface to the most commonly used features of PHREEQC. The more advanced features (Inverse Modeling, Kinetics, and Advection/Transport) are not included in this option. To utilize these features, you may load one of the USGS's full versions of graphical interfaces as explained below.

PHREEQC (Advanced)

This will load an external window, containing one of the USGS's PHREEQC interfaces (PHREEQC-Interactive or PHREEQC for Windows). This will allow you to use the full features of the PHREEQC modeling program, using your AquaChem samples as initial solutions.

NOTE: You must have one of the mentioned external graphical interfaces installed and configured in the PHREEQC section of the preferences in order to use this feature.

Generate PHT3D Input

This option allows you to create PHT3D input files for use with PHT3D in Visual MODFLOW versions 4.2 or higher. For more information on this option, please refer to “Generate PHT3D Input” on page 366.

3.9 Window Menu

Using this menu you can arrange the windows in AquaChem master window. Below the **Arrange Icons** option there is a list of all open windows, so you can easily navigate between them without shrinking them to arrange on a common screen.

Tile Vertical

Arranges open windows one above the other.

Tile Horizontal

Arranges open windows side by side.

Cascade

Cascades open windows.

Arrange Icons

Organizes icons of minimized windows along the bottom of the parent window.

3.10 Help Menu

Contents

Displays the AquaChem On-line Help.

Index

Displays the Help index.

About

Displays the AquaChem version number and information on how to contact Schlumberger Water Services.

This concludes a review of the AquaChem menu items. The following sections describes the components of an AquaChem database.

4

Plots

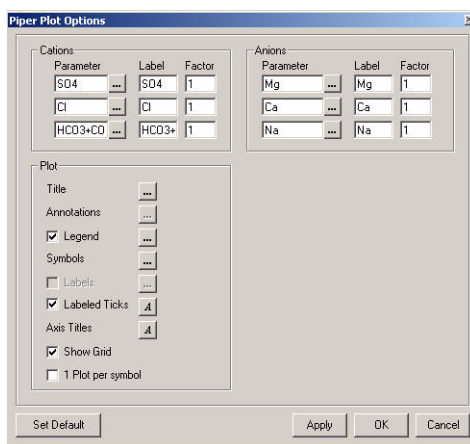
When you select **Plots** from the main menu and then **New**, a sub menu will appear with a list of the twenty three (29) available plot types:

- Box and Whisker (Grouped)
- Box and Whisker (Multiple Parameters)
- Box and Whisker (Multiple Stations)
- Box and Whisker (Time)
- Depth Profile
- Detection summary
- Durov Plot
- Geothermometer Plot
- Giggenbach Triangle
- Histogram
- Ludwig-Langelier Plot
- Map Plot
- Meteoric Water Line (MWL) Plot
- Pie Plot
- Piper Plot
- Probability
- Quantile Plot
- Radial Plot
- Scatter Plot
- Schoeller Plot
- Stacked Bars (Distance)
- Stacked Bars (Stations)
- Stiff Plot
- Ternary Plot
- Time Series (Multiple Parameters) (Multiple Parameters)
- Time Series (Multiple Parameters) (Multiple Stations)
- Time Series (Stacked Bars)
- Time Series (Statistics Summary)
- Wilcox Plot

AquaChem allows you to create multiple plots for the same data set and view these plots simultaneously within the Windows environment. Each of these plots is explained in greater detail later in this chapter. The following section describes some of the features that are common to all plots.

4.1 Common Plot Features

Although each graph type has unique characteristics, there are also many graphical features and options that are common for each one. When you select any of the graph types to plot, a **Plot Options** dialogue similar to the one below, will appear with default settings for the necessary parameters and plot settings.



After the plot has been created, there are two ways to access the **Plot Options** dialogue:

- Click **View>Options** from the main menu when a plot window is the active window; or
- **Right-mouse** click on the **centre** of the plot window.

The **Plot Options** dialogue is typically divided into two major parts: the **Parameter/ Series Options** and **Plot Options**.

The Parameter/Series Options portion of the dialogue allows you to set up various parameters of the plot, while the Plot Options portion allows you to set up such plot features as the legend, symbols, title, and labels. Simply click the ... button to load the appropriate dialogue.

- **Parameters Dialogue** - Contains information on the parameters/series used in the plot, axis labels and font options, axis titles and ranges, and toggles to turn the plot gridlines or axis ticks on/off.
- **Title Dialogue** - Contains options for plot title, position, and font size.
- **Symbols Dialogue** - Contains options for symbols used in the plot, proportional symbol options, etc.
- **Legend Dialogue** - Contains options for displaying a plot legend, legend title, and display features.
- **Labels Dialogue** - Contains options for displaying and customizing symbol labels.
- **Format (Axis) Dialogue** - Contains options for customizing the axis values and appearance.
- **Annotations Dialogue** - Contains the options for adding one or more text annotations onto your plot.

Each **Options** dialogue has the following buttons located at the bottom of the dialogue:

The **[Apply]** button will apply the current plot settings to the selected plot type.


The **[OK]** button will apply the current plot settings to the selected plot type and will close the plot options dialogue.

The **[Cancel]** button closes the plot options dialogue.

The **[Set Default]** button will save the current plot settings as defaults for that plot type. This includes plot parameters, grid lines, interval, axes titles, legend appearance and symbol settings, but does not include plot titles. The default settings will be applied every time you create a **NEW** plot of that type, with the current database.

The **Axes/Parameters** dialogue has several buttons and functions which are common for most (but not all) plot types:



The add button allows you to add new parameters to the plot. The add button will load the parameter list dialogue, and allow you to select a new parameter from the list of available parameters in your database. Simply choose the desired parameter, press the **[Select]** button and this parameter will be included in the plot. When you have chosen all needed parameters, click **[Close]** to return to the options. The  button beside a required parameter performs the same function.



The delete button will remove the selected parameter from the plot parameters list.



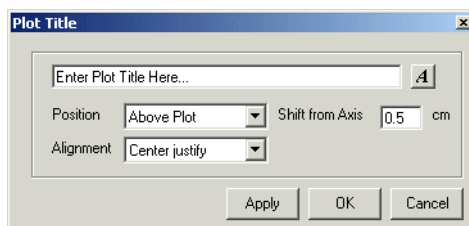
The up-arrow button allows you to shift the selected parameter upwards in the list of parameters.




The down-arrow button allows you to shift the selected parameter downwards in the list of parameters.

4.1.1 Title Dialogue

Clicking on the  button beside **Title** launches the following dialogue:



In this window, you can enter a **Plot Title** and edit the title font by pressing the  button. In addition, you can change the **Position**, and **Alignment** of the plot title.

Plot Title

Defines the title of the selected graph type. Enter the plot title in the text field provided. If no text is entered, then no title will be plotted.

Position

Defines the position of the plot title as either **Above Plot** or **Below Plot**.

Alignment

Defines the alignment of the plot title as either **Left justify**, **Centre justify**, or **Right justify** in the plot window.

Shift From Axis

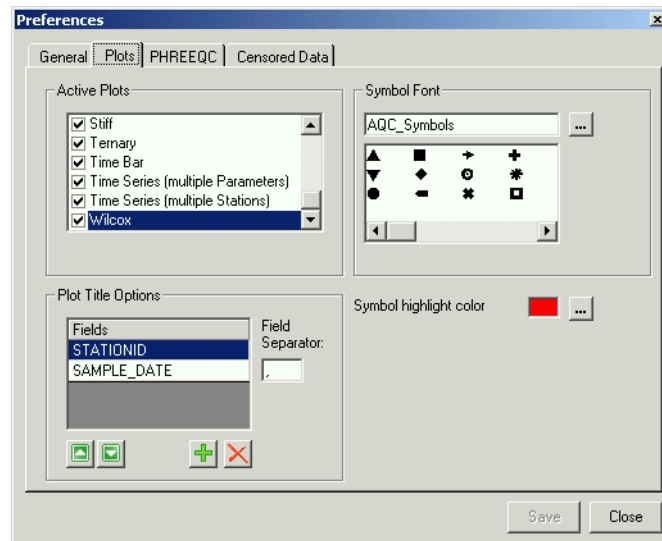
Defines the vertical distance of the plot title above or below the plot.

Automatic Plot Title Option

AquaChem includes a useful utility which will allow you to automatically assign plot titles based on any station or sample description parameter, or a combination of these parameters. This is very practical especially when you are creating a large number of plots.

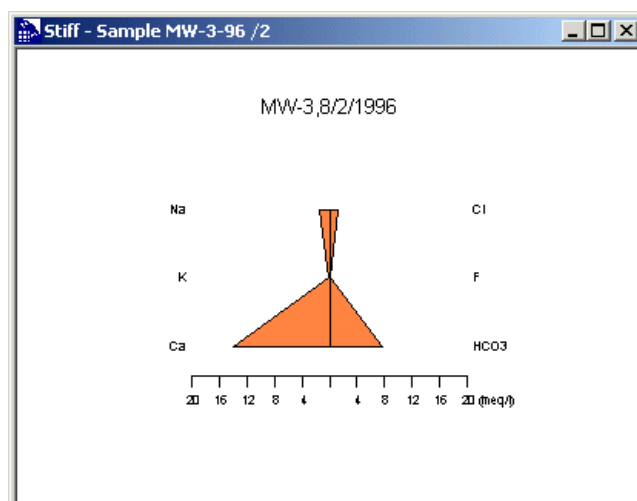
NOTE: This feature is applicable only to plot types that represent one distinct sample per plot (**Pie**, **Radial** and **Stiff** plots only).

To setup the Automatic Plot Title features, close all plot windows and ensure the sample list is the active window. Select **File** from the main menu, then **Preferences**. From this dialogue, choose the **Plots** tab and a window similar to the one shown below will appear.



Under the **Plot Title Options** section (in the lower-left corner), you can specify which parameters you would like to appear when using the automatic title option. Parameters can be arranged, and added/removed using the up, down, add, or remove buttons. Once you are satisfied with the parameters, you must specify a **Field Separator**. This will be the character that separates the parameters in the title of the plot. You can use any character as the field separator, however the most common are a hyphen, comma, or semi-colon.

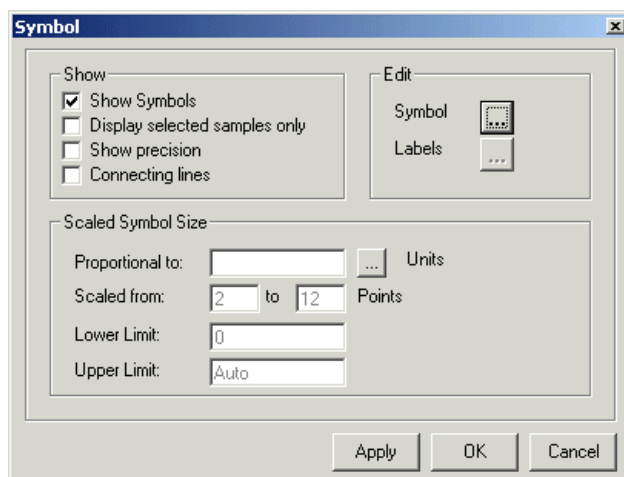
Below you can see an example of the plot that uses the automatic titling feature with specified parameters:



NOTE: By default, all **Pie**, **Radial**, and **Stiff** plots will be titled automatically with the fields specified in the **Preferences** dialogue. To change the plot title after the plot has been created, simply type in a new plot title in the **Plot Title** field, and press **[OK]**.

4.1.2 Symbols Dialogue

The options in the **Symbols** dialogue allow you to edit the symbol appearance settings.




Show frame

In this frame you find the following options:


- **Show Symbol** - show/hide symbols using this check-box
- **Display selected samples only** - display only the samples selected from the Samples List using this check-box (need to pre-select the samples)
- **Show precision** - show/hide precision bars using this check-box (Precision value must be entered for this particular parameter in the Sample Details window)
- **Connecting lines** - show/hide connecting lines using this check-box

Edit frame

Symbols

Click the  button to open the **Define Symbol or Line** dialogue. In this dialogue, you can define the names and status of the symbol groups, the shape and color of the symbols, and the line-type and color of lines that appear in the plots (refer to the Define Symbol or Line section in Chapter 3 for a description of the symbol groups settings).

Labels

Click the  button to open the **Labels** dialogue. In this dialogue you can specify the labels for each individual data point or for multiple data points at once. Select the text, font, and position for the label. For more detail see **Labels dialogue** section later in this chapter.

Scaled Symbol Size frame

Proportional to

This option allows you to scale the size of the plot symbols based on a user-defined parameter (TDS, pH, Cond, etc.). This allows you to represent an additional parameter on your plot. For example, in a scatter plot, you can display information on three parameters, instead of only two (X and Y). Once the parameter is selected, you can choose different units for this parameter.

Scaled from ... to ... Points

Defines the minimum and maximum radius of the scaled symbol size. The minimum symbol size will be plotted for the lower limit of the proportional parameter value, while the maximum symbol size will be plotted for the upper limit of the proportional parameter value. The minimum radius ensures that all symbols will be visible even if the proportional parameter value is zero.

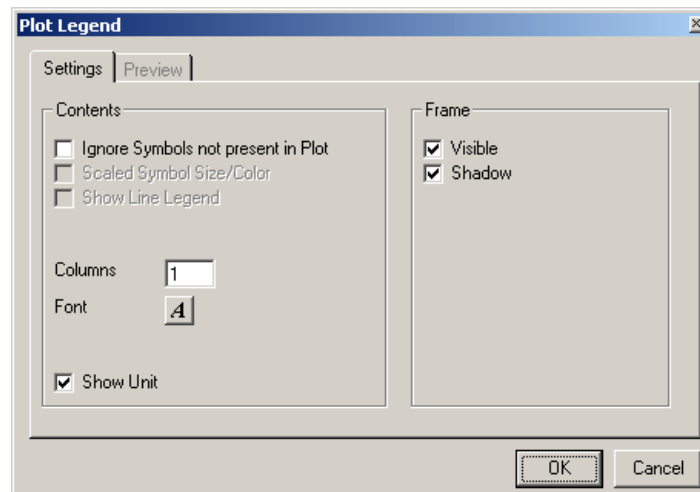
Lower and Upper Limit


Defines the minimum/maximum value of the proportional parameter to use for plotting a proportional symbol size. The lower limit is the value below which the symbol size does not get any smaller; i.e. all samples with a proportional parameter value less than the lower limit, will use the smallest symbol point size; likewise for the upper limit. The purpose of the lower limit is simply to set a value below which the symbol sizes do not change.

If you want to exclude symbols from the plot based on a specified criteria, then you should create a query to filter out the undesirable samples.

4.1.3 Legend dialogue


Most plots have legend options, as shown below:



You can access these options either by clicking the  button beside **Legend** in the **Plot Options** dialogue, or by right clicking on the plot legend itself.

Contents frame

Symbol Names

Show/hide symbol names using this check-box. When the legend is visible, the **Symbol Names** from the currently selected symbol group will be displayed on the plot. Use the font button  below the options to edit the font of the symbol names as they appear in the legend.

Ignore symbols not present in plot

When this option is active, the plot will show only those symbols which are used in the current plot. For example if 10 symbols are active, but the current selection of samples only uses 2 of them, AquaChem will ignore those symbols that are not present in the plot and display just the 2 symbols in the legend.

Scaled Symbol Size/Color

This option is only active if you have symbol sizes plotted proportional to a parameter value, under the **Symbols** options. When this is activated, the legend will show a scale for the proportional symbol sizes.

Show Line Legend

Show/hide line legend using this check-box.

Columns

Specify the number of columns for displaying legend items

Frame frame

Visible

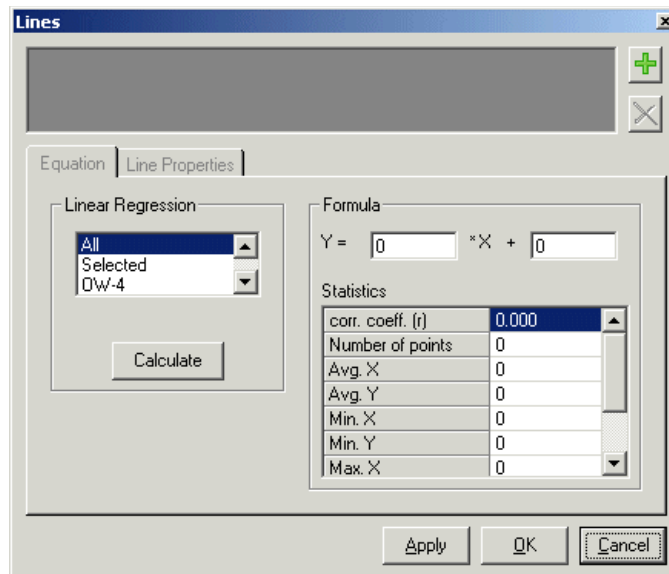
Show/hide box around the legend using this check-box.


Shadow

Show/hide the legend shadow effect using this check-box.

4.1.4 Line Dialogue

AquaChem allows you to put a calculated or user-defined regression line on your plot. For Time Series and Box and Whisker plots horizontal lines can also be added to represent line features such as Maximum Concentration Limit.



To create a new line click the  in the upper right corner. The defined line follows the basic equation of $y=mx+b$ and once a new line is created you need to specify these constants for the line. There are two ways of doing so: calculate a linear regression trend or enter user defined values.

Equation tab

Linear Regression frame

This frame allows you to calculate the linear regression constants based on various selections:

- **All** - calculates linear regression using all available stations
- **Selected** - calculates linear regression using only selected series
- **Individual stations (names depend on the database)** - calculates linear regression based on one station

Once you've selected the data series, click the **[Calculate]** button to calculate the coefficients.

NOTE: The selection must include more than one point.

Formula frame

This frame allows you to enter custom values for the linear regression equation constants. If you used the **Linear Regression** frame to calculate values based on data

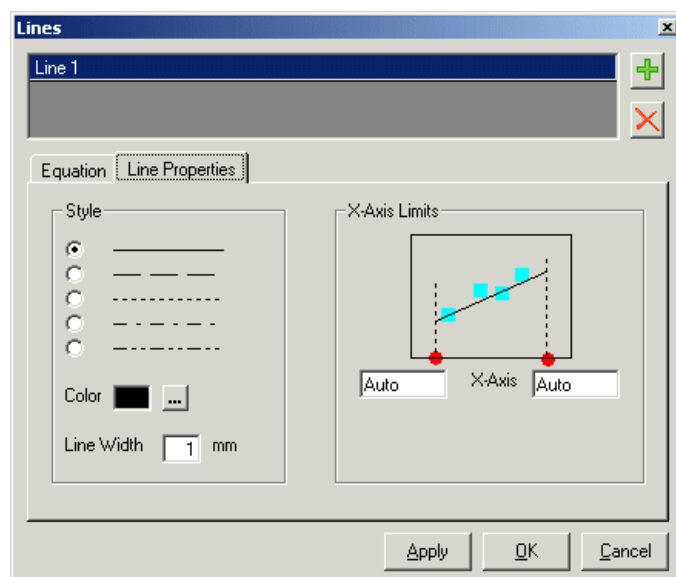
points, these values will be displayed in this frame. If the **Calculate** feature was used, this frame will also display statistics of the calculations.

Adding a Horizontal Line (Time Series and Box and Whisker plots)

To add a horizontal line to a Time Series or Box and Whisker plot to represent a feature such as a Maximum Concentration Limit line, simply create a new line in the **Lines** dialog and edit the line name as you would like it to appear in the plot legend. Then in the **Formula frame**, enter the Y-axis value where you would like the line to be positioned in the ***b*** field of the $y=mx+b$ formula. The line can be included in the plot legend using the Legend dialogue (See section 4.1.3 for more information on the Legend dialogue).

Line Properties tab

Use this tab to customize the line appearance on the plot.



Style frame

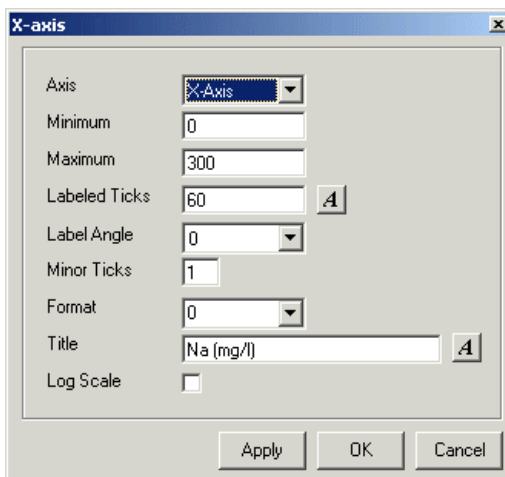
Set the **Style**, **Color**, and **Width** of the line.

X-Axis Limits frame

Set the maximum and minimum X-value for the generated line. If left and right are set to **Auto**, the line will span the entire plot.

4.1.5 Format dialogue

This dialogue is only available in plots on X-Y style plots with linear axes.



The image shows a software window titled "X-axis" with a close button in the top right corner. Inside the window, there are several input fields and buttons. The "Axis" field is a dropdown menu showing "X-Axis". The "Minimum" field contains the value "0". The "Maximum" field contains the value "300". The "Labeled Ticks" field contains the value "60" and has a small button with the letter "A" next to it. The "Label Angle" field is a dropdown menu showing "0". The "Minor Ticks" field contains the value "1". The "Format" field is a dropdown menu showing "0". The "Title" field contains the text "Na (mg/l)" and has a small button with the letter "A" next to it. The "Log Scale" field is a checkbox that is currently unchecked. At the bottom of the window, there are three buttons: "Apply", "OK", and "Cancel".

Axis

This combo box allows you to switch between the available axes.

Minimum/Maximum

Allows you to customize the minimum and maximum axes values. AquaChem will determine a default minimum/maximum value based on all of the data plotted on the graph. If you enter a new value and you wish to return to automatic calculation, enter "A" in this field.

Labelled Ticks

Defines the interval value of the labelled ticks for the corresponding axis. AquaChem will determine a default value for the labelled ticks such that the plot will have 5 labelled ticks on the corresponding axis. If you enter a new value and you wish to return to automatic calculation, enter a value of "A" in this field.


Label Angle

Allows you to specify the angle to the axis at which the labels appear. Tilted tick labels are recommended for long labels which run the risk of overlapping on smaller plots.


Minor Ticks

Defines the number of minor ticks between each of the labelled ticks. The default value is 1. Only integer values are accepted.

Format

Sets the numeric format for the labels on the axes (e.g. Auto, 0, 0.0, 0.00, 0.0E+00, etc.). Beside **Format**, click  button in the combo box to select from the list of available numeric formats.

Title

Defines the title of the corresponding axis. If a title field is left empty, a default title equal to the internal name of the parameter will be selected by AquaChem. If you want to omit one or both axes titles, type a space in the text field(s). Use the  button to change the font.

Log Scale

Some plots allow you to show the plot data using a log scale for either the X or Y axis, or both. This allows you to normalize the data set, if there is a significant range in the data.

4.1.6 Labels dialogue

Use this dialogue to set up the label display properties. This dialog is only available once the plot is created. All changes made to the labels are automatically updated in the plot.

ID	SAMPLEID	WATERTYPE	SAMPLE_DATE	STATIONID
1	MW-1-92	Ca-CO3-SO4	8/15/1992	MW-1
2	MW-1-93	Ca-Na-SO4-Cl-HCO3	6/1/1993	MW-1
3	MW-1-94	Ca-Na-SO4-Cl	6/15/1994	MW-1
4	MW-1-95	Ca-Na-SO4-Cl-HCO3	7/30/1995	MW-1
5	MW-1-96	Ca-Na-SO4-Cl-HCO3	7/28/1996	MW-1

Parameter

☒ Assign parameter result to all samples
☐ Assign parameter result to selected samples only

none

Edit Label

Label

☐ Visible
☐ Connecting line to Symbol

Font

Position

Close

In the grid at the top of the dialogue, select the samples for which you want to display a label. Click and drag the mouse pointer to select the stations in a continuous list, or hold down the **Ctrl** key to select individual samples.

When the **Labels** dialogue is loaded, any samples that are currently selected on the plot (or in the Active List) will be highlighted for you.

Parameter frame

If you wish the labels to be displayed only for the selected samples, click the radio button beside **Assign parameter result to selected samples only**. Otherwise, leave the default setting of **Assign parameter result to all samples**.

Use the combo box to select the parameter that will be used as the label. As soon as you do so, the labels will be displayed for selected (or all) samples.

Edit Label frame

You may use this frame to customize a particular label, or create a custom label for any sample. To do so, select a sample from the grid at the top of the dialogue. If this sample already has a parameter label, it will be displayed in the **Label** field. If this sample has no label assigned to it, the **Label** field will be blank.

Label

The label text is normally filled automatically from a database field. You may, however, edit this text or even enter all labels manually.

Visible

Use this check-box to display/hide the specified label for the selected sample(s).

Connecting line to Symbol

Use this check-box to display/hide a line connecting the symbol with the label. The connecting line is useful for densely placed symbols.

Font

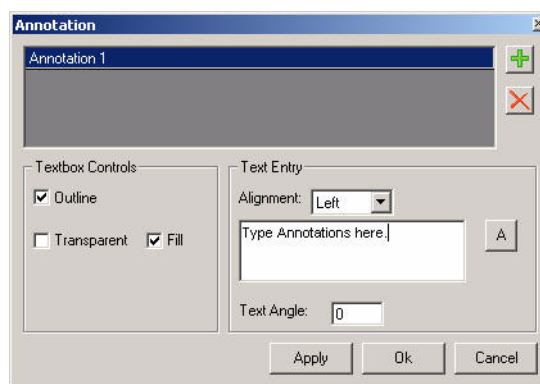
Click the **A** button to change the label font.

Position frame

Use the arrow buttons to position the labels. The label position is updated as soon as you click the arrow button.

4.1.7 Annotations Dialogue

This dialogue provides the options for adding text annotations onto your plots. This dialog is only available once the plot is created.



The green plus will add an annotation while the red x will delete an annotation. There are also options for an outline for the annotation box or whether to have the box transparent or filled white.

Under the Text Entry portion of the dialogue you will find options for the alignment, font and font angle. The text is entered into the white box.

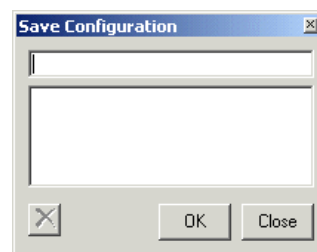
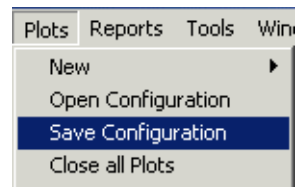
4.2 Plot Configurations


Once you have created the desired plot, or combination of several plots, the **Save Configuration** option under the **Plots** menu is enabled. This allows you to save the current configuration of open plots and their settings, to your database.

The configuration includes axis settings, selected samples, symbol settings, etc.

When you select **Save configuration**, the dialogue shown on the right side will appear prompting you for a name for the Plot Configuration. Simply type in a name for the plot configuration.

Once you are finished, click **[OK]**.

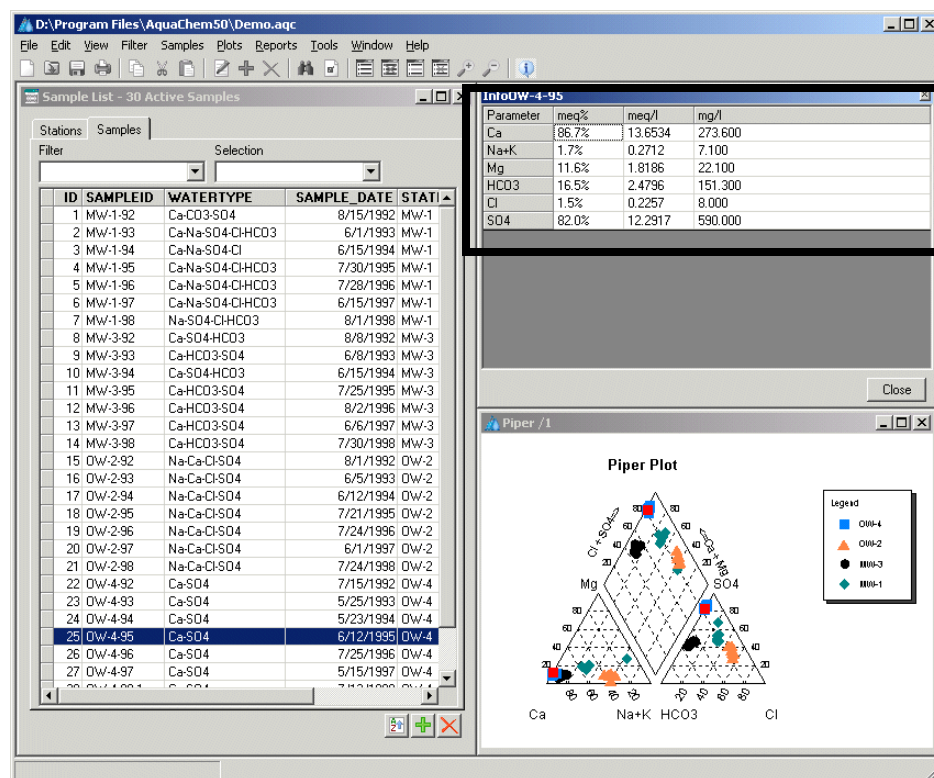


To recall a saved plot configuration in the future, use the **Plots > Open Configuration** option. Simply select the plot configuration you desire, then click **[OK]**. This will load the plot(s) and their settings. When a plot configuration is loaded, the sample selection used for this plot will also be reset in the Active List window. Aquachem allows for multiple plots configurations to be open concurrently. To remove any unwanted or outdated plot configurations, simply select the respective item from the list, and press the  button. Once you are finished, click **[Close]**.

4.3 Save, Show, and Identify Plot Data

Show Sample Data

If this option is selected from the **Plots** menu, AquaChem will display a small dialogue with the data for the selected sample. Simply click on a sample point from the plot and the **Info** data dialogue will appear. Each time you click on a new sample point, the new data is loaded into dialogue. An example is shown below:



NOTE: You may need to re-arrange the positions of your windows in order to see the Sample Data dialogue.

Identify Plot Data

The **Identify** option allows you to link the plotted data to the samples in the active list. This allows you to click on a point in the graph and view the corresponding sample in the active list. This is useful when you have a large number of samples displayed on one plot, and you want to identify outliers.

NOTE: You may specify the color in which the symbol is highlighted on the plot in the **Plots** option under **File>Preferences** described in chapter 3.

There are three options available for selecting and identifying points on a plot: **None**, **Selected Plot**, and **All Plots**. The default mode to identify points is on All Plots; this means that all data points on all plots are linked to the database. These three options are described below:

None

If this option is selected, then no symbols will be highlighted on the plot when the samples are selected. The data points on the graphs are not linked to the database which means that samples are not identified. When you have several open graphs and a large database with many samples, it may be convenient to have no link between the graphs and the database in order to increase the speed of the various AquaChem operations.

Selected Plot

Only the data points in the active graph are linked to the database. The chosen point is activated (highlighted) and selected in the sample list. This mode is useful if you are working with a large data set and want to focus on just one graph.

All plots

The sampling point of the corresponding sample is activated on all plot windows and on the sample list. If you are working with several graphs, you can identify a sample in all plots as well as in the sample list by clicking near a point in any of the plots, or by clicking on a sample in the active list. The sample will become selected in the active list, and the corresponding data points will be highlighted in red in all the open plot windows.

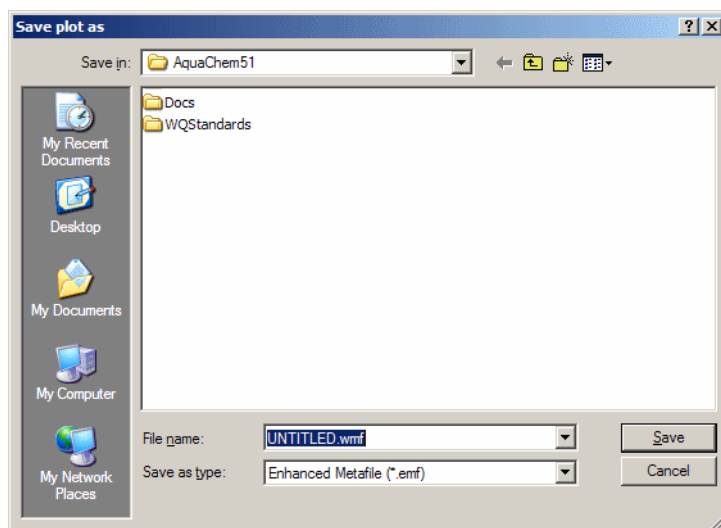
4.4 Printing and Exporting Plots

Once you are satisfied with your plot(s) and their design, you have three options for producing output:

- The plot can be saved as a Windows Metafile (.WMF) and printed from an external application, or inserted into a report;
- The plot can be copied to the Windows clipboard and pasted into another supported application (ex. a graphics program, MSPaint, or a word processor);
- The plot can be printed as is, or can be incorporated into a Printing Template which can include project information, company details, and your company logo.

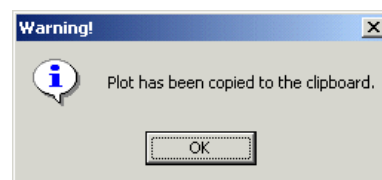
4.4.1 Export as Metafile

When you have any plot window open, you can save the plot as a Windows Metafile (graphics format) by selecting **File** from the main menu and then **Export>Image**. In the **Save Plot As...** dialogue that appears, enter a filename for this file and press **[Save]**. You can now open and manipulate this image using a graphics editor or insert this file into a report. When using this feature, each plot window must be saved individually as a metafile.




4.4.2 Copy Plot to Clipboard

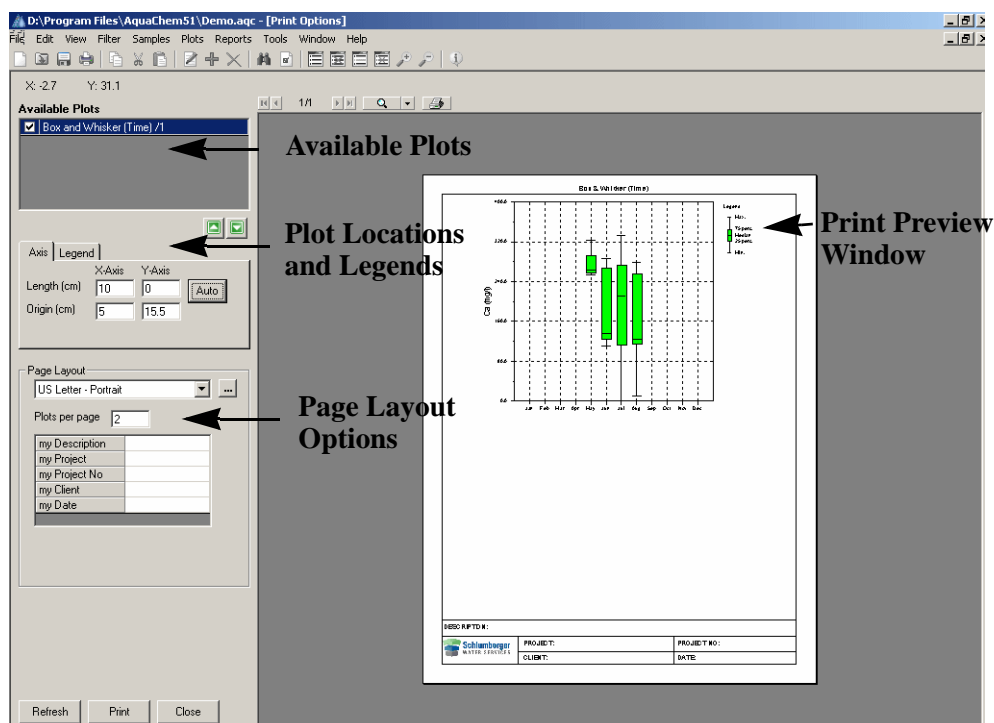
To copy the selected plot to the windows clipboard, select **Edit** from the main menu and then **Copy**. If successful, a confirmation dialogue will appear.



The plot can then be pasted directly into another application such as a word processor or a graphics editor.

4.4.3 Printing

To print one or more of the on-screen plots, select **File** from the main menu and then **Print**, while the plots window is active. Alternatively, you can click the  icon from the AquaChem toolbar. A **Print Options** window will appear, as shown in the figure below:



The **Print Options** window has the following components:

Available Plots: Contains the list of plots that were open when **Print** option was selected, and thus are available for printing.

Plot Legends and Locations: Lists the X and Y location of the plot and the legend (if available).

Page Layout Options: Lists the available plot templates and options for customizing the report.

Print Preview Window: Provides a preview of the printed page.

To print one or more plots, you must select the desired plot(s) from the list of the **Available Plots** by placing a check mark in the box beside the appropriate plot. The selected plot(s) will appear in the **Print Preview** window and will be automatically sized and arranged to fit the page.

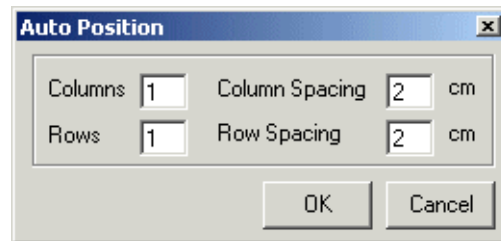
To select a plot, click once on the plot list to activate it, and then click once in the box beside the plot name. A check mark will be added to the box and this plot should appear in the **Print Preview** window. To load additional plot(s), simply click once with your mouse in the box beside the plot name(s).

To remove a plot from the **Print Preview** window, simply click once in the box to remove the check mark beside the appropriate plot.

Arranging the Plots

The order of the plots can be easily modified using the arrow buttons below the list of plots. Selected plots can be moved up or down using these buttons. The position of each plot on the page can be modified in the **Axis** tab by entering a new X, Y origin; the size of each plot can be modified by entering a new X, Y length.

Clicking the **[Auto]** button will load the following dialogue:



In this dialogue edit the grid used to optimally position plots on the page.

If needed, click the **[Refresh]** button at the bottom of the screen to update the print preview.

Plots per page option allows you to spread the plots over several pages, as well as specify the number of plots to be arranged on each page.

Print Preview Window

The **Print Preview** window provides a “What You See Is What You Get” (WYSIWYG) preview of the printed page. The page magnification value can be adjusted using the magnifying glass above the preview window. This will enlarge or shrink the appearance of the page preview on your screen. The page coordinates (cm) for the mouse pointer location are located in the upper left corner of this window.

Selecting a Print Template

The **Page Layout** frame allows you to select from a list of pre-defined print templates which contain fields for descriptions of the plot(s), the project, client, and company information. In this frame there is also a **Plots per page** field. This allows the user to specify how many plots they would like to appear on each page when printing multiple plots.

Plot arrangement and page layouts can be saved as templates at anytime by selecting **Save as Template** from the **File** menu.

AquaChem includes the following pre-defined print templates:

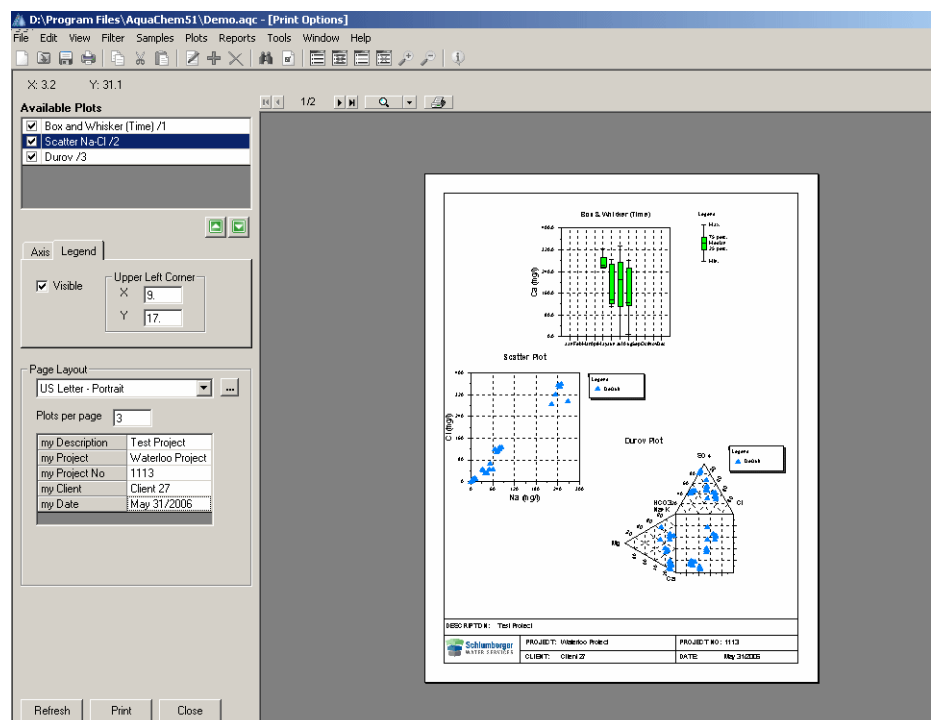
- US Letter - Portrait
- US Letter - Landscape
- A4 - Portrait
- A4 - Landscape

The default template selection is "**None**", meaning no print template is selected. If "**None**" is selected you have the option to enter a two line title at the top of the page plus a one line footer at the bottom of the page.


If the pre-defined print templates are not satisfactory for your needs, you can easily create your own customized print templates using the **Template Designer** option available by clicking the '...' button beside the template selection list. For a detailed description of this component, please refer to the Template Designer section in chapter 3.

If you select one of the pre-defined print templates, a list of available descriptor fields will appear and the **Print Preview** window will be updated to reflect the layout of the selected template. Fill in the project specific plot description fields under the **Page Layout** options. Depending on the template, you may enter information for: CLIENT, PROJECT, PROJ #, DATE, DESCRIPTION.

An example of the **Print Preview** window with Fields is shown below:



Once you have entered the descriptive information for the plot, press the **[Refresh]** button to refresh the print preview.

When you are satisfied with the print layout, simply click the **[Print]** button in the lower left corner of the window, select **File > Print** from the main menu, or click the  icon.

Alternatively, you may select **Edit > Copy** from the main menu, and the plot, along with the print template, can be copied to the clipboard, and pasted into an external application.

To return to the main AquaChem window, click the **[Close]** button.

4.5 Plot Details

This section describes each of the AquaChem plots and the options available for each plot.

4.5.1 Box and Whisker

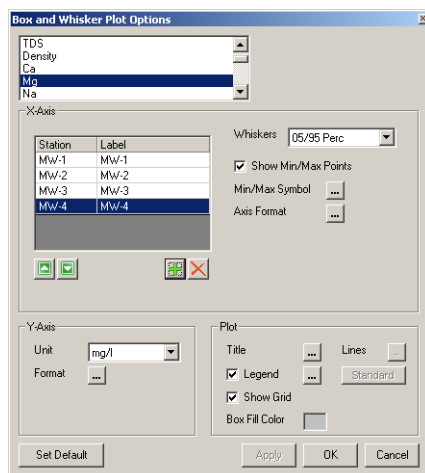
The **Box and Whisker** plot displays a statistical summary of any measured database parameter(s). It is composed by a central box showing the spread of the bulk of the data (interquantile range) and the whiskers, showing the length of the tails.

The Box and Whisker plot can be displayed in three forms:

- **Box and Whisker (Grouped)**
- **Box and Whisker (Multiple Parameters)**
- **Box and Whisker (Multiple Stations)**
- **Box and Whisker (Time)**

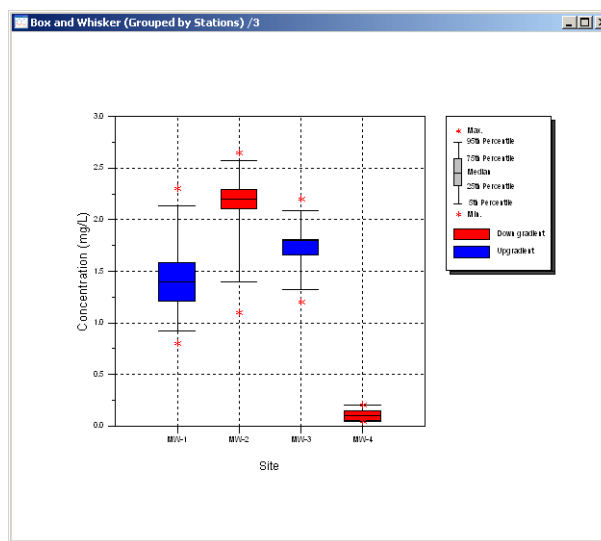
X-Axis frame

When the **Box and Whisker (Grouped)** plot is used, samples are grouped by sample name, and each box is colored according to the current symbol configuration. This plot can be useful for comparing station box and whisker plots belonging to different domains/areas, *e.g.*, upgradient/downgradient.



First, select the desired parameter from the top list of parameters, and then add one or more stations to the stations list. When the station box and whisker plots are created, they will be colored based on the current station symbol configuration. For more information on assigning station symbols, see “Assign Symbol” on page 155.

An example of a **Box and Whisker (Grouped)** is shown below:



If you select **Box and Whisker (Multiple Parameters)** plot then select one or more stations group in the **Series** field, and then specify one or more parameters under the **Parameters** list. The options for this plot are shown below.

Box and Whisker Plot Options

Series: Mw-1, Mw-2, Mw-3, Mw-4

X-Axis

Parameters	Label
Na	Na
Ca	Ca
Mg	Mg
Cl	Cl
HCO3	HCO3
SO4	SO4

Whiskers: 05/95 Perc

☒ Show Min/Max Points

Min/Max Symbol: ...

Axis Format: ...

Y-Axis

Unit: mg/L

Format: ...

Plot

Title: ... Lines: ...

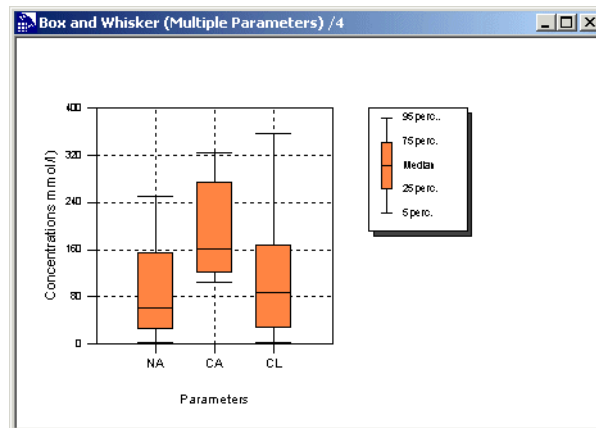
☒ Legend

☒ Show Grid

Box Fill Color: [Blue]

Buttons: Set Default, Apply, OK, Cancel

An example of the **Box and Whisker (Multiple Parameters)** plot is shown in the figure below.



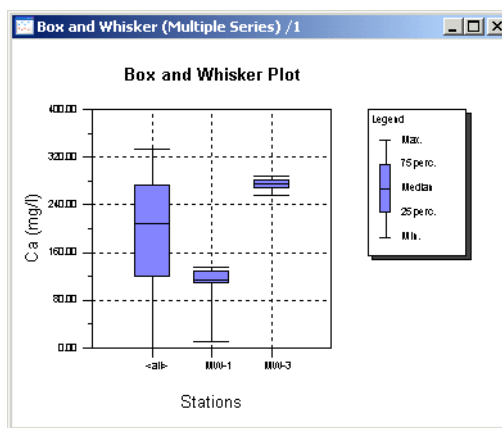
If you select **Box and Whisker (Multiple Stations)** plot then you will see the **Parameters** and **Series** fields become reversed. You can highlight one or more parameters in the **Parameter** field and then enter one or more stations in the **Station** list. The options for this plot are shown below.

The figure shows the "Box and Whisker Plot Options" dialog box. It contains the following sections:

- Sample Depth:** A list box containing "Sample_Depth", "Temp", "pH_field", "pH_lab", and "Eh".
- X-Axis:**
 - A table with "Station" and "Label" columns:

Station	Label
MW-1	MW-1
MW-2	MW-2
MW-3	MW-3
MW-4	MW-4
 - Whiskers: A dropdown menu set to "05/95 Perc".
 - ☒ Show Min/Max Points
 - Min/Max Symbol: A button with three dots.
 - Axis Format: A button with three dots.
- Y-Axis:**
 - Unit: A dropdown menu set to "mg/L".
 - Format: A button with three dots.
- Plot:**
 - Title: A button with three dots.
 - Lines: A button with three dots.
 - ☒ Legend: A button with three dots.
 - ☒ Show Grid
 - Box Fill Color: A color selection button showing a blue square.

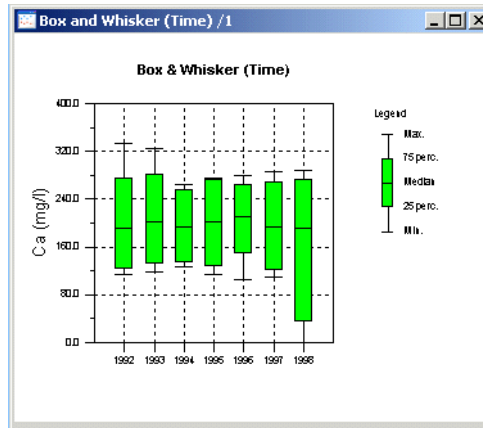
An example of the Box and Whisker (Multiple Stations) plot is show in the figure below:



If you select **Box and Whisker (Time)**, the analysis will be performed on all samples in the Sample List for one or more parameters. AquaChem can aggregate the samples by month, season or year. Note that the season and month group may include samples from different years and are particularly useful for seasonal analysis.

The figure shows the "Box and Whisker Plot Options" dialog box. The "Sample" list includes "Depth", "Temp", "pH field", "pH lab", and "Eh". The "X-Axis" section has "Aggregate by" set to "Year" and "Whiskers" set to "05/95 Perc". The "Y-Axis" section has "Unit" set to "mg/L" and "Format" set to "...". The "Plot" section has "Title" set to "...", "Legend" checked, "Show Grid" checked, and "Box Fill Color" set to "Standard".

An example of the **Box and Whisker (Time)** plot is shown in the figure below:



Select the statistics to be displayed as whiskers using the **Whiskers** combo box.

Whereas the extent of the central box is standardized (25%, 75% quantiles), the extent of the whiskers is not. The whiskers combo box, therefore, lets you choose, which position of the tails of the underlying population should be used to displayed by the whiskers: Min/Max extent, 5/95% or 10/90% quantiles. If the whiskers are defined as a quantile value, then the minimum and maximum values can be added as a further information to the plot and will be shown as points above and below the whiskers.

Click the button beside **Format** to produce the **Axis** dialogue. Use this dialogue to customize the x-axis appearance.

Y-Axis frame

Unless all parameters you wish to plot are unitless, this frame allows you to select the display unit using the **Unit** combo box.

Click the button beside **Format** to load the **Axis** dialogue. Use this dialogue to customize the y-axis appearance and apply log transformation if necessary.

Plot frame

Click the button beside **Title** to load the **Title** dialogue.

Check the box beside **Legend** to enable the plot legend. Click the button to load the **Legend** dialogue and customize the display options.


Check the box beside **Show Grid** to display the gridlines.


Click the button beside **Lines** to load the **Lines** dialogue. In this plot the **Line** option may be used to draw a horizontal line representing the standard for the or one of

the plotted parameters. This can be done manually by creating a line $y=mx+q$ where $m=0$ and q equal to the standard for this parameter.

If the standard value is defined in the active water standard, then clicking on the **[Standard]** button will automatically create this line for you.

Fill Color/Pattern

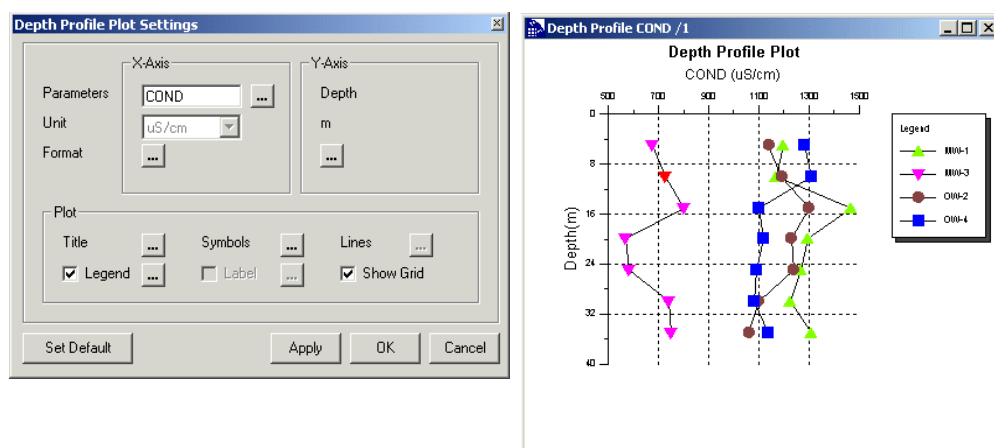
Click the  button beside **Color** to select the fill color for the box plot.

Click the  button beside **Pattern** to choose the display pattern for the box plot.

For more details on this plot, please refer to the article: Tukey J.W., 1977: *Exploratory Data Analysis*, Addison-Wesley, Reading, Massachusetts, USA. 1977, pp. 39-43.


4.5.2 Depth Profile

The **Depth Profile** plot displays the change in a parameter value as sample depth changes. An example of a **Depth Profile** plot and the corresponding **Depth Profile Plot Settings** dialogue is shown in the figure below.




The **Depth Profile** plot is similar to a borehole log. The plot displays the change of a measured parameter over a measured sampling depth. In order to create this plot, you must have an entry for the parameter Sample_Depth for one or more samples in your database. You may then plot the **Depth** (on the Y-Axis) against any measured parameter (on the X-Axis).


X-Axis and Y-Axis frames

Click the  button beside the **Parameter** field in the **X-Axis** frame to select the parameter to be plotted on the x-axis. The parameter for the y-axis frame is **Depth** and cannot be changed.

Select the units for the x-axis parameter using the **Unit** combo box in the **X-Axis** frame. The unit for the y-axis depends on the **Depth** parameter unit in the database and cannot be changed in this dialogue.

Click the  button beside **Format** in the **X-Axis** and **Y-Axis** frames to access their respective settings dialogues.

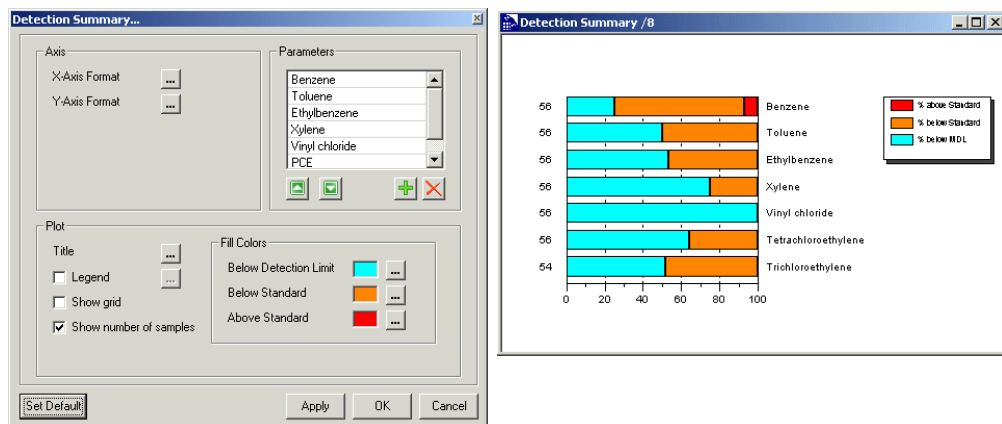
Plot frame

Click the  button beside **Title**, **Symbols**, **Lines**, **Legend**, and **Labels** to access their respective settings dialogues. The **Legend** and **Labels** features must be activated (the box beside them must be checked) before their dialogues can be loaded.

Check the box beside **Show Grid** to display the gridlines.

4.5.3 Detection summary

The **Detection Summary** plot is used to visualize the relative proportions of samples above the specified water quality standards. The plot shows a summary bar for every parameter specifying the percentage of values where the contaminant has not been detected, where it has been detected but is below the water quality standard and the percentage of samples exceeding the water quality standard. Below is the **Detection Summary** plot options and the corresponding plot:







On the left side of each detection summary bar you can see the number of samples that have a value for the parameter that this bar represents. The parameter name is listed on the right side of each detection summary bar. The X-axis lists cumulative percent values from 0 to 100. You can calculate the number of samples that exceed a particular standard (or are non-detects) by multiplying the percent “length” of the particular colored segment by the number of samples used to produce a particular detection summary bar.

NOTE: Parameter values that contain “<” symbol will be included in “below MDL” category.


Axis frame

Click the  button beside the **X-axis Format** and **Y-axis Format** to load their respective options dialogues.

Parameters frame

Use this frame to select the parameters for which you wish to produce the detection summaries. Use the  button to add parameters (select from the presented list), the  button to delete parameters, and  and  buttons to change the order in which they appear on the plot.


Plot frame

Click the  button beside **Title** and **Legend** to access their respective dialogues. The **Legend** feature must be activated (checked) before the options dialogue can be loaded.

Use the check box beside **Show Grid** to show/hide the gridlines.

Use the check box beside **Show number of samples** to show/hide the number of samples displayed to the left of the detection summary plot.

Fill colors frame

Use this frame to customize the display colors for detection summary plot. Use the  button to access the color selection dialogue.

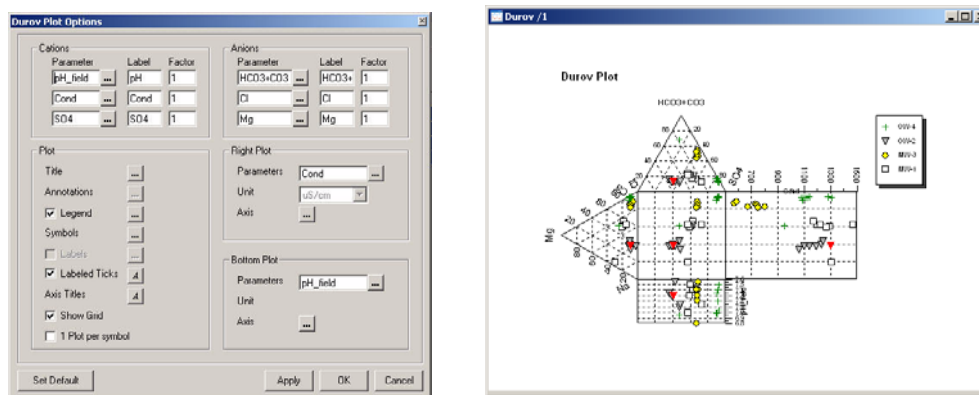
4.5.4 Durov Plot

The trilinear **Durov** plot is based on the percentage of major ion milliequivalents. The **Cations** and **Anions** values are plotted on two separate triangular plots and the data

points are projected onto a square grid at the base of each triangle. In addition, the Durov plot allows for the direct comparison of two other groundwater parameters, typically pH and the total dissolved solids (TDS). The **Durov** plot is an alternative to the Piper plot which is described later in this chapter.

Since the data points are projected along the base of the triangle, which lies perpendicular to the third axis in each triangle, information about the concentration of the vertex element (the third element) is lost in the square grid. Changing the orientation of parameters in both triangles may improve your ability to detect distinct groups.

An example of **Durov** plot and the corresponding **Durov Plot Options** dialogue is shown in the figure below.




Cations and Anions frames

The **Cations** specify the parameters of the left triangle. Default settings are the major cations **Na**, **Ca**, and **Mg**, however any other parameter can be selected (e.g. gas composition, trace elements etc.). The **Anions** generally specify the parameters for the upper triangle. Default settings are the major anions **Cl**, **SO4**, and **HCO3**, however any other parameters can be selected. Note that measured Alkalinity is a valid anion parameter, provided the respective parameter is known by the system as being the Alkalinity. This may be done on the **File>Database>Alias** tab by assigning the respective parameter to the Measured Alkalinity. If using the alkalinity in the anion triangle, change the respective displayed label field to $\text{HCO}_3 + \text{CO}_3$.

Data from poorly concentrated parameters can be multiplied by a common factor to prevent data point accumulation on a base line when plotting parameters with very different concentrations. Type the multiplication **Factor** in the field beside the parameter.


Both the Cations and Anions frames also contain a **Label** field for each parameter. This allows customization of the parameter labels on the plot diagram. For example the parameter 'Ca' can be edited in the **Label** field to displayed as 'Calcium'.

Plot frame

Click the  button beside **Title**, **Symbols**, **Legend**, and **Labels** to access their respective options dialogues.


The **Legend** and **Labels** features must be activated (checked) before the options dialogues can be loaded.

Check the boxes beside **Labeled Ticks** and **Show Grid** to display the tick values and grid lines on the plot.


Click the  button to change the font of the **Labeled Ticks** and the **Axis Titles**.

Click the 1 Plot per Symbol option to create one Durov plot for each station.

Right Plot frame

The right plot is located to the right of the middle grid of the Durov plot. Choose which parameter to show on this plot by selecting the  button located beside the **Parameters** field. You may also choose the desired **Unit**, and customize the appearance of the **Axis**.

Bottom Plot

The bottom plot is located below the middle grid of the Durov plot. Choose which parameter to show on this plot by selecting the  button located beside the **Parameters** field. You may also choose the desired **Unit**, and customize the appearance of the **Axis**.

4.5.5 Geothermometer Plot

Geothermometer plots can be used to test the quality of geothermometer estimates for a given geological and hydrogeological situation. Different geothermometers can produce very different results when applied to the same water sample(s). Each geothermometer has a set of conditions which must exist for reasonable values to be calculated. For example, if the in-situ temperature was not sufficient to attain rock-water equilibrium, the aquifer composition may be different from the one assumed by the geothermometer. Using conventional geothermometer graphs (log K-1000/T plots), you can check the quality of the calculated temperature if you have in-situ temperature

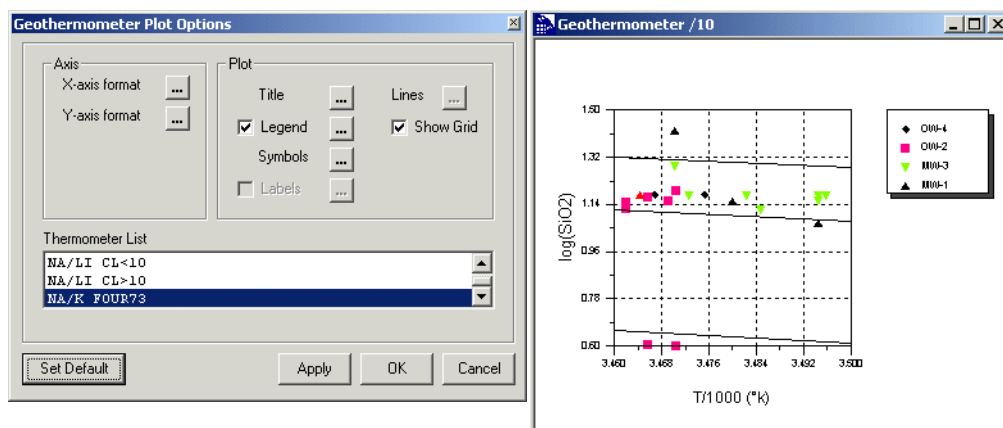
measurements. You should collect all analyses from the literature for which in-situ temperatures have been measured. Also note the aquifer lithology of the samples. When you start a geothermal study, search for references to samples which compare closely with the aquifer of the water you are studying, and check which geothermometer gives the most reliable estimates. This thermometer is likely to give the most reliable results for the water you are studying.

Geothermometer plots can be used to:

- Check the applicability of a thermometer on a set of samples.
- Search for the geothermometer for which you observe the best fit.
- Develop new chemical thermometers for parameters or parameter ratios which show linear behavior in this plot.

For a list of the available Geothermometers, please see the Thermometers options, in the **File > Database** dialogue.

The **Geothermometer** plots display most commonly used geothermometers. An example of a **Geothermometer** plot and the corresponding **Geothermometer Plot Options** dialogue is shown in the figure below.




The following section describes some of the features and options of the **Geothermometer Plot** that are not covered in the Common Plot Features section.

Geothermometers are generally expressed by the formula:


$$t(K^{\circ}) = \frac{1000 \cdot a}{b + \log(x)}$$

where, x is the SiO_2 concentration or Na/K , etc., $\log(x)$ values plot on a straight line as a function of $1000/T$. The coefficients a and b are obtained empirically from deep borehole data with in situ temperature measurements.

Axis frame

Click the  button beside **X-axis format** and **Y-axis format** to access their respective options dialogues.

Plot frame

Click the  button beside **Title**, **Legend**, **Symbols**, **Labels**, and **Lines** to access their respective options dialogues. The **Legend** and **Labels** features must be activated (checked) before the dialogues can be loaded.

Use the **Show Grid** check box to show/hide the plot grid lines.

Thermometer List

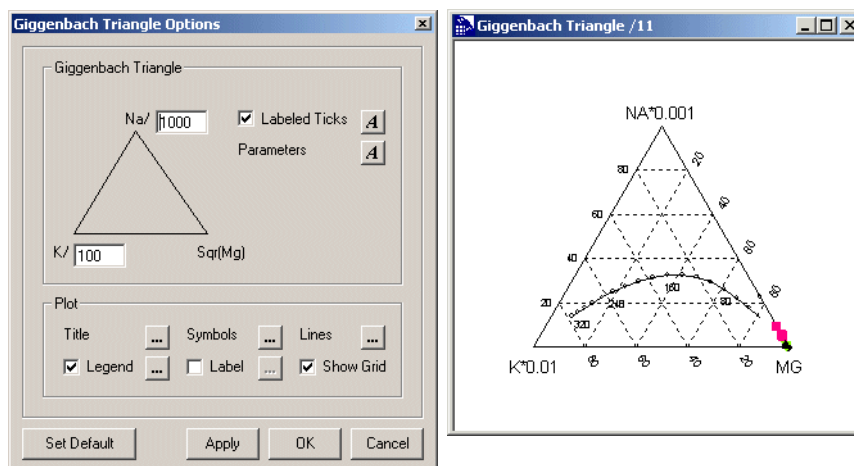
Select (highlight) the thermometer to be used for the plot.

For more details on Geothermometers, please refer to the following article:

Kharaka et al. 1989.

4.5.6 Giggenbach Triangle

The Giggenbach triangle provides a visual aid to determine the water-rock equilibrium. An example of the **Giggenbach Triangle** plot and the corresponding **Giggenbach Triangle Options** dialogue is shown in the figure below.



The Giggenbach Triangle (K-Mg-Na Triangle) representation allows you to verify the extent to which water-rock equilibrium has been attained.

The triangle is comprised of three zones:

- Immature waters (at the base);
- Partially equilibrated waters (in the middle); and
- Fully equilibrated waters (along the curve).

Depending on where the composition of a given sample lies within this triangle, you can estimate the extent of rock-water equilibrium. For mature waters falling near the upper curve, you can estimate the temperature as well.


Giggenbach Triangle frame

Enter the division factors for **Na** and **K** in the text fields provided. The default values for the plot are Na/1000 and K/100.

Use the **Labeled Ticks** check box to show/hide the tick values.

Click the  button to modify font for the tick values and parameter labels.

Plot frame

Click the  button beside **Title**, **Symbols**, **Legend**, and **Labels** to access their respective options dialogues. The **Legend** and **Labels** features must be activated (checked) before the options dialogues can be loaded.

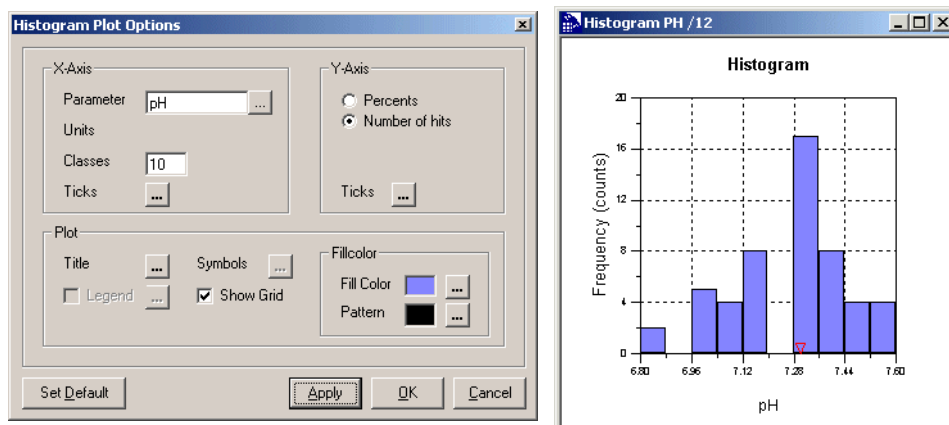
Use the **Show Grid** check box to show/hide the plot grid lines.

For more details on the Giggenbach Triangle, please refer to the following article:

Giggenbach, W.F., 1988. *Geothermal solute equilibria*. Derivation of Na-K-Mg-Ca geoindicators. *Geochim Cosmochim. Acta*, 52: pp. 2749-2765.


4.5.7 Histogram

Two of the oldest methods for summarizing data distributions are the frequency and histogram plots. Both the histogram and the frequency plot use the same basic principles to display the data: dividing the data range into units, counting the number of points within the units and displaying the data as the height within a bar graph. An example of a **Histogram** plot and the corresponding **Histogram Plot Options** dialogue is shown in the figure below.



You can easily identify a single sample in a multi-sample histogram plot by clicking on the sample in the Active Samples List. A red triangle will be displayed on the histogram, indicating the range interval of the selected sample (this is a unique feature of AquaChem and is not normally found in other histogram plotting software). Clicking on a bar on the graph will select all the samples in the Active List which are included within the interval range defined by the histogram bar you selected.

X-Axis frame

Use the  button beside **Parameter** to select parameter to be plotted on the histogram. This field can also handle parameter ratios, sums, or differences (e.g. Na/Cl, Ca+Mg, Cl-SO₄). Enter these parameter combinations into the parameter field manually. Unless the parameter you've chosen is unitless (e.g. pH), the **Unit** combo box will allow you to select the units for the parameter.

The **Classes** field defines the number of data classes (or intervals) into which the range of parameter values will be subdivided. Each data class will be represented by a single bar on the histogram. Type the number of classes (bars in the histogram) in the provided field.


Click the  button beside **Format** to load the corresponding axis options dialogue.

Y-Axis frame


The number of samples within each data class is referred to as the frequency of occurrences. This can be expressed as either a percentage of the total number of samples (**Percents**), or as the number of samples within each data class (**Number of hits**). Select the desired option.

Click the  button beside **Format** to load the corresponding axis options dialogue.

Plot frame

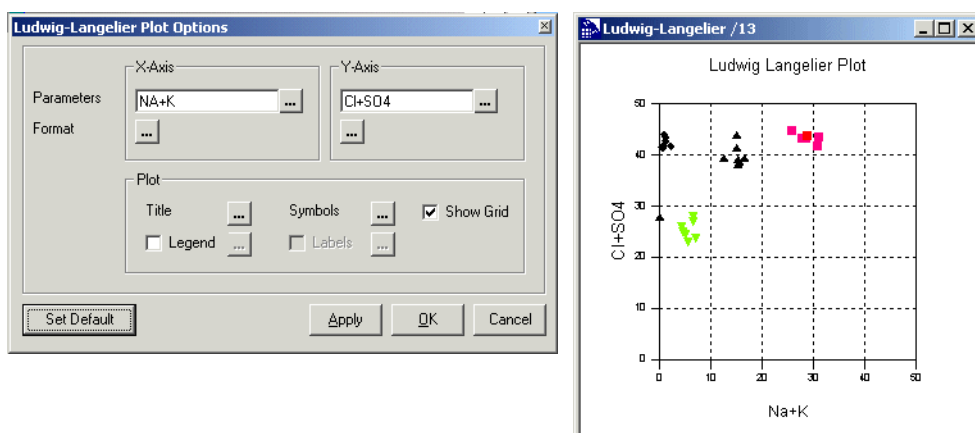
Click the  button beside **Title** to load the title options dialogue. Check the box beside **Show Grid** to display the grid lines.

Fill Color and Pattern frame

The color and pattern of the bars in the graph can be modified using the options beside **Color** and **Pattern**. Simply press the  button to access the various options.

4.5.8 Ludwig-Langelier Plot

The **Ludwig-Langelier** plot allows you to quickly see patterns and correlations between the major cations and anions for multiple samples. An example of the **Ludwig-Langelier** plot and the corresponding **Ludwig-Langelier Plot Options** dialogue is shown in the figure below.



The **Ludwig-Langelier** square plot is similar to the projection areas of the **Piper** and **Durov** plots. By convention, the sums of selected cations are plotted on the **X-Axis** and the sums of selected anions are plotted on the **Y-Axis**. Each axis ranges from 0 to 50 meq%. Sample points are calculated as follows:

$$\Sigma_{anions} = Cl + SO_4 + HCO_3$$

$$\Sigma_{cations} = Ca + Mg + Na + K$$

$$\% Na = 50 \frac{Na}{\Sigma cat}$$

$$\% Cl = 50 \frac{Cl}{\Sigma an}$$


Suitable groupings of cations and anions are selected and plotted as percentages. Generally, this type of graph is used to plot %Na+%K against %HCO₃+%SO₄. In this plot, %Ca+%Mg and %Cl are also fixed:


$$\%Ca + \%Mg = 50 - (\%Na + \%K)$$

$$\%Cl = 50 - (\%HCO_3 + \%SO_4)$$


All major elements can be displayed in one plot with the **Ludwig-Langelier** plot, however like the **Piper** and **Durov** plots, the plot displays relative ratios rather than absolute concentrations.

X-Axis and Y-Axis frames

Click the  button beside the **Parameter** field to select the first of the parameters. The remaining parameters must be added manually.

Click the  button beside **Format** to load the axis options dialogue for the corresponding axis.

Plot frame

Click the  button beside **Title**, **Symbols**, **Legend**, and **Labels** to load the corresponding dialogue. The **Legend** and **Labels** features must be activated (checked) before their options dialogues can be accessed.

Use the **Show Grid** check box to show/hide the plot grid lines.

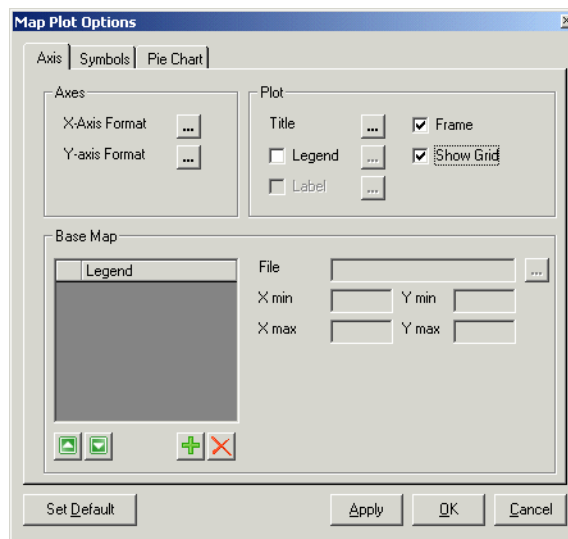
4.5.9 Map Plot

The **Map** plot reads the X, Y coordinates for the station location, and displays this on a blank grid or on a site map of your site. The **Map** plot can import and overlay an AutoCAD (.DXF) file, an ESRI shape (.SHP) file, or a common graphics (.BMP, .JPG, and .GIF) file as a basemap of the site or study region defining the major physical and geographical boundaries and geological characteristics.

The **Map** plots can be used to simply display the station locations throughout the study region, or to interpret spatial trends in the physical or chemical characteristics of each sample using proportional symbol sizes, or by plotting **Pie**, **Radial** or **Stiff** graphs at each sample location. The Radial and Stiff symbols may be exported to ESRI polygon shapefile format.

NOTE: You must have at least one station in your database with X and Y co-ordinates in order to create a **Map** plot. Also, station coordinates must be expressed in a projected coordinate system, and X and Y values must be positive.

When you choose a **Map** as the plot type, the following **Map Plot Options** dialogue will appear:




Axis tab

Axes frame

Click the  button beside **X-axis format** and **Y-axis format** to access the corresponding dialogues.


Plot frame

Click the  button beside **Title**, **Legend**, and **Label** to access their respective options dialogues. The **Legend** and **Labels** features must be activated (checked) before the options dialogue can be loaded.

Use the **Frame** and **Show Grid** check boxes to show/hide the frame around the map and the plot grid lines.

Base Map frame

AquaChem allows you to load multiple maps in various file formats as the base map for the plot. You can load any of the following types of files: .DXF, .SHP, .BMP, .JPG, and .GIF. If the format of the file you wish to load is not listed above, use a graphics editor to convert it into one of the listed formats.

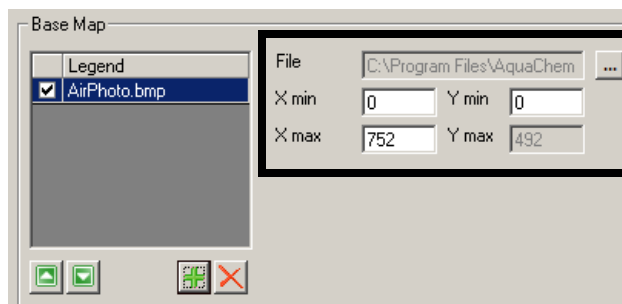
To add a basemap to your plot click the  button. In the **Base Map File** dialogue that is loaded select the appropriate file type and navigate to the desired file. Click **[Open]** to load the map and the file name will appear in the **Base Map** frame as shown below.

If you loaded a .DXF or a .SHP file, the **min X/max X** and **min Y/max Y** coordinates are entered automatically. For the ESRI shape files you will also have the options of changing the line width and color.


For shape files, additional options will appear allowing you to specify line width and color.



NOTE: The images must be georeferenced in feet or meters (the units have to be the same as the station X-Y coordinates). AquaChem does not support Lat/Long coordinate system.

If you load a common graphics file (.BMP, .JPG, or .GIF), the program automatically assigns one meter (or foot) per pixel in the picture. This image has to be manually georeferenced.



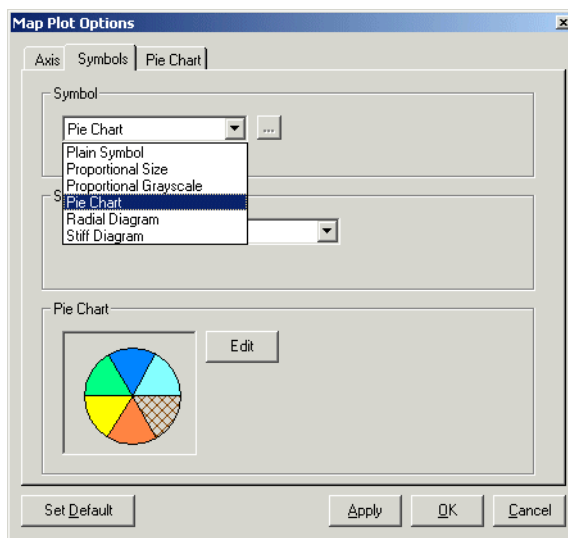
Simply enter the new coordinates for 2 corner points of the map.

If you wish to change the source file for the currently highlighted map, click the  button to load a new file.

You may load multiple maps into your map plot. Use the  and  buttons to change the order in which they appear in the plot and the check box beside the name to show/hide the respective map image.

Symbols tab

The **Symbols** tab in the **Map Plot Options** dialogue allows you to choose a symbol to represent the samples on the map plot




Symbol frame

There are six symbol types to choose from:

- Plain Symbol
- Proportional Size
- Proportional Greyscale
- Pie Chart
- Radial Diagram
- Stiff Diagram


Depending on the type of symbol you choose, the lower preview frame will show different plot and the third tab (available only for **Pie Chart**, **Radial Diagram**, and **Stiff Diagram**) will present different options to customize the chart.

Plain Symbol

The **Plain Symbol** type will plot the symbol shapes, sizes and colors as they are defined for each sample group. Press the  button beside the field to access the **Define Symbol or Line** dialogue for viewing or modifying the symbol settings for the database.


Proportional Size Symbols

The **Proportional Size** symbol type will plot the defined symbols at each sample location with a symbol size determined by the value of a selected parameter. The sample values for the proportional parameter will be used to determine the radius of the symbol at each sample location (e.g. samples containing higher values of the proportional parameter will have a larger symbol radius).

To change the proportional parameter, type in the internal name of the desired parameter in the **Parameter** field, or click the  button and select another parameter from the list.

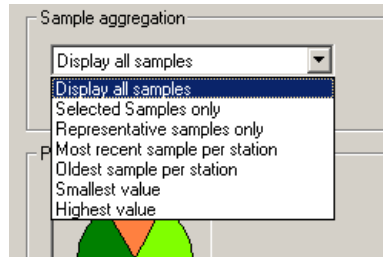
The **Radius** field is used to set the minimum and maximum symbol radius (in points). The minimum symbol radius corresponds to the **Lower Limit** value of the proportional parameter, while the maximum radius corresponds to the **Upper Limit** value. Use a non-zero minimum radius to plot symbols at sample locations containing very low values of the proportional parameter. The symbol radius at each sample location is determined using a linear interpolation between the **Upper Limit** value and the **Lower Limit** value.

Proportional Greyscale

This symbol type will display a **Greyscale** symbol at each sample location, with the scale determined by the value of a selected parameter. The sample values for the proportional parameter will be used to determine the greyscale color of the symbol at each sample location (e.g. samples containing higher values of the proportional parameter will have a darker greyscale). To change the proportional parameter, simply type in the Internal name of the desired parameter in the **Parameter** field, or click the  button and select another parameter from the parameters list. Sample with values less than the Lower Limit value are assigned a white symbol color, while points with values equal to or greater than the Upper Limit value are assigned a black color fill.

Sample Aggregation

If you wish to display only one sample for every station in your project, this frame allows you to select the criteria according to which this sample will be chosen:



Select the appropriate criterion. Select **None** if you wish all samples to be considered.

The following aggregation options are available:

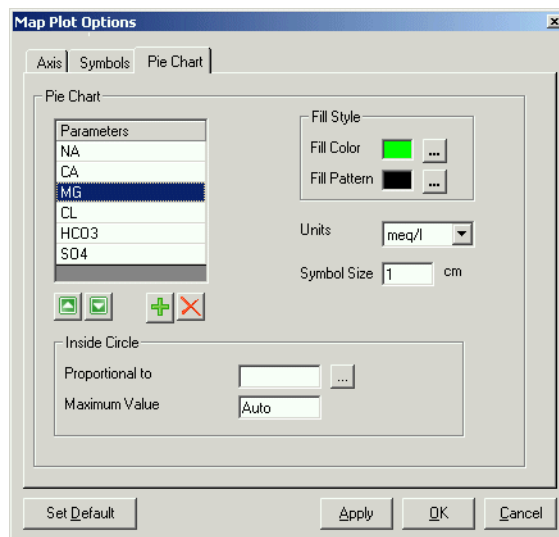
- **Selected Samples only** - only the samples currently selected in the Samples List will be displayed
- **Representative samples only** - only samples that are marked as Representative in the Samples List will be displayed
- **Most recent sample per station** - one sample with the latest date will be displayed for each station
- **Oldest sample per station** - one sample with the earliest date will be displayed for each station
- **Smallest value** - presents an option to choose a parameter based on the value of which one sample (with the smallest value) will be displayed
- **Highest value** - presents an option to choose a parameter based on the value of which one sample (with the largest value) will be displayed
- **Closest to average** - presents an option to choose a parameter based on the value of which one sample (with the value closest to the station mean) will be displayed

Preview frame





This frame is available only for **Pie**, **Radial** and **Stiff Diagrams**. It presents a generalized view of the chosen plot and the **[Edit]** button. Clicking on this button will move you to the third tab which, depending on the plot type you chose, will be one of the described below.

Pie Chart tab


This symbol type will display a **Pie** chart of the selected parameters at each sample location.



Parameters list

Click the  button to add parameters to the pie chart and the  to delete them. Use the  and  buttons to change the order of the parameters.


Fill Style frame

Use the  buttons beside **Fill Color** and **Fill Pattern** to specify these features.

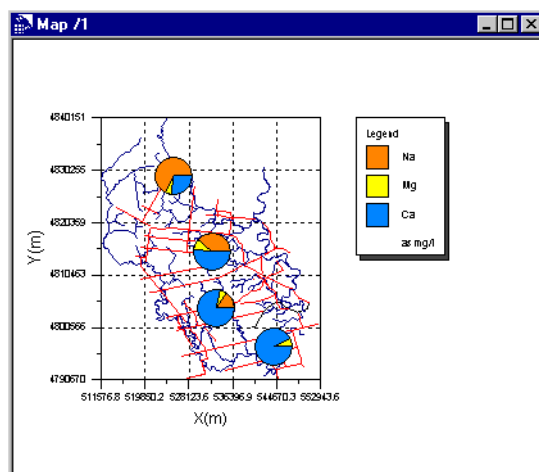
Use the **Units** combo box to select the units for the parameters that require them.

Type in the symbol size in centimeters in the **Symbol Size** text box.

Inside Circle frame

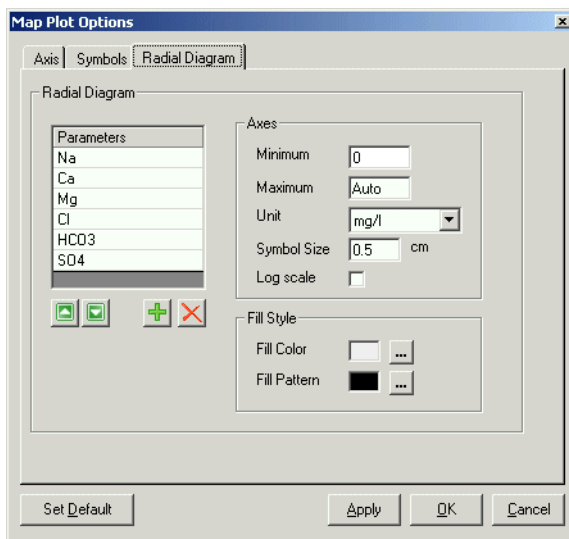
If you wish, you can display one parameter as an inside circle of the pie plot. Click the  button to select the parameter from the list and enter the **Maximum Value** for the parameter. The samples containing a **Proportional** parameter concentration equal or greater than the **Maximum Value** will have an inside circle with the maximum radius. The inside circles for samples with **Proportional** parameter concentrations less than the **Maximum Value** will have a proportionally smaller radius.

An example of a **Map Plot** with **Pie** charts as sample symbols is shown below.







Radial Diagram tab

This plot will display a Radial Diagram of the selected parameters at each sample location.



Parameters list

Click the  button to add parameters to the radial diagram and the  to delete them. Use the  and  buttons to change the order of the parameters.

Axes frame


Set the **Minimum** and **Maximum** values for the axes. AquaChem will select the optimum values, however they may not be what you want.

Select the units for the parameters using the **Units** combo box.

Set the **Symbol Size**. This value is the radius of the symbols as they appear on the printed output. Depending on your monitor's resolution they may seem different on the screen.

You may log-transform your data using the **Log scale** check box.

Fill Style frame

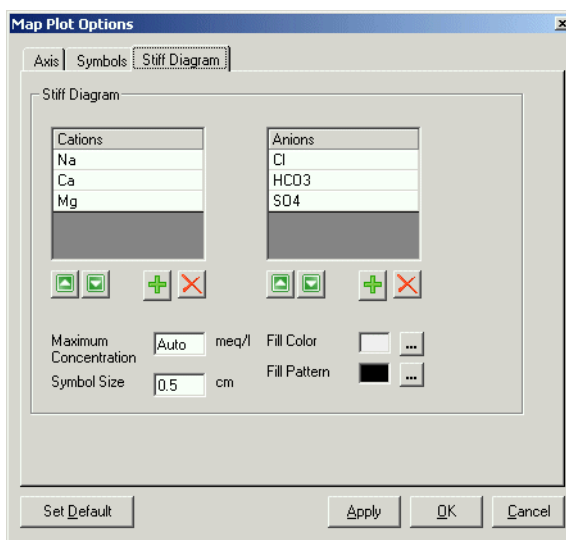
Use the  buttons beside **Fill Color** and **Fill Pattern** to specify these features.

Use the **Units** combo box to select the units for the parameters that require them.





Type in the symbol size in centimeters in the **Symbol Size** text box.

Stiff Diagram tab

This plot will display a **Stiff** diagram of the selected parameters at each sample location.




Cations and Anions list

A **Stiff diagram** plots an equal number of cations and anions. Click the  button to add parameters to the **Cation** and **Anion** list and the  to delete them. Use the  and  buttons to change the order of the parameters.

NOTE: You must have an equal number of **Cations** and **Anions** which is greater than zero to generate this plot.

The **Maximum Concentration** values will represent the uniform maximum value for each parameter. The minimum axis value is always zero.

The **Symbol Size** field contains the setting for the size of the Stiff diagrams plotted at each sample location.

The **Fill Style** (**Fill Color** and **Fill Pattern**) for each parameter can be modified by clicking on the target parameter, then choosing a new Color or the Pattern by clicking on  button beside **Fill Color** and **Fill Pattern** fields.

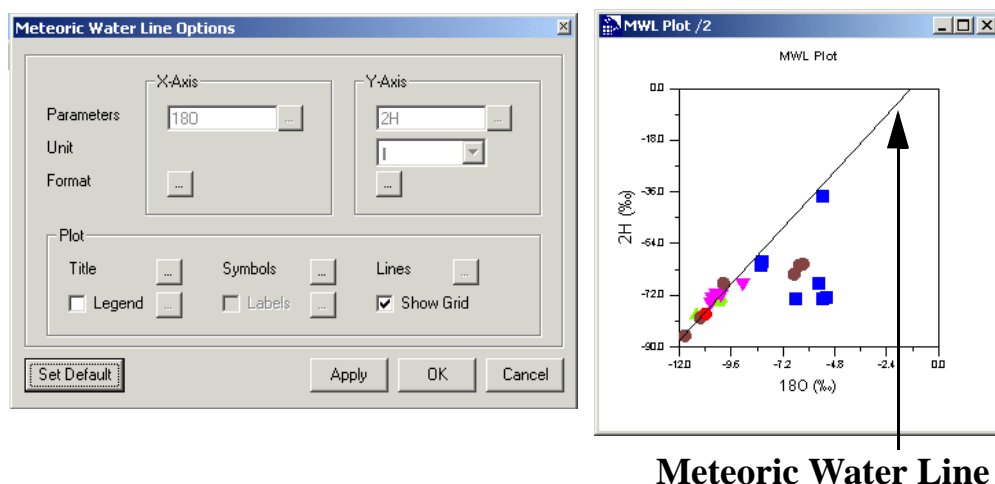
Once you have configured all options to your satisfaction, click **[OK]** to generate the plot. You can always access the options dialogue by right-clicking on the plot.

4.5.10 Meteoric Water Line (MWL) Plot

The **Meteoric Water Line** plot is the standard way to portray measurements of stable isotopes from water (^{18}O and ^2H). It consists of a scatter plot showing ^{18}O on the X-axis and ^2H on the y-axis. The meteoric water line is defined by the following equation:


$$^2\text{H} = 8.2 \times ^{18}\text{O} + 10.8$$

Most precipitations and groundwaters fall closely on this line. In cases where the measurements are shifted from this line, the type of shifting provides important information regarding the process leading to the observed shift (e.g. rock water interaction, evaporation, mixing with seawater etc.).




Meteoric Water Line

X-Axis and Y-Axis frames

The **Parameters** and **Units** are selected automatically for this plot and cannot be changed. You can change the axis options by clicking on the  button and loading the corresponding **Axis** dialogue.

Plot frame

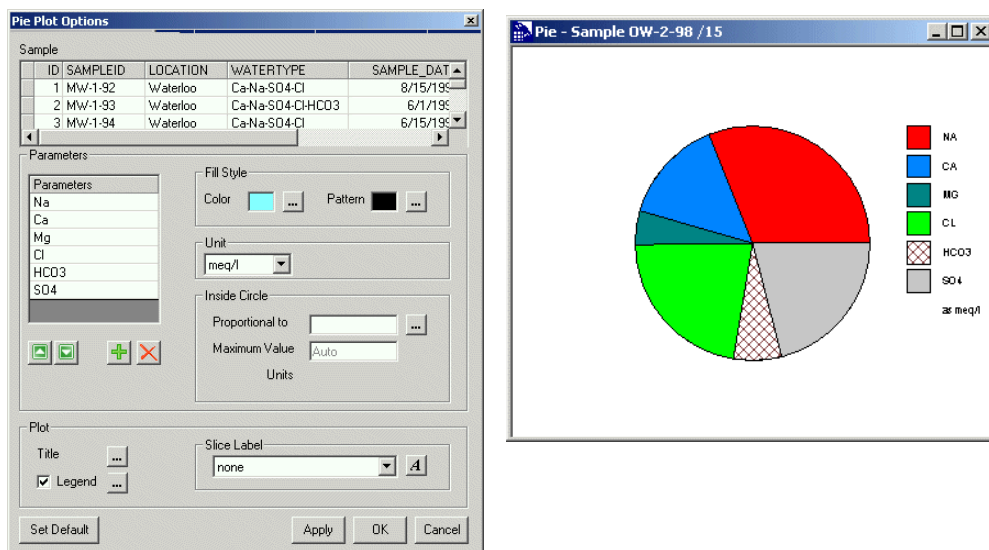
Click the  button beside **Title**, **Symbol**, **Lines**, **Legend** and **Labels** to access the corresponding options dialogues. The **Legend** and **Labels** features must be activated (checked) before the dialogues can be loaded.

Use the **Show Grid** check box to show/hide the plot grid lines.

NOTE: this plot is only active if your database contains the parameters 18O and 2H and if these parameters are mapped in the **Alias** section.

4.5.11 Pie Plot

The **Pie** plot is a simple way of showing parameter portions in a sample. An example of the **Pie** plot and the corresponding **Pie Plot Options** dialogue is shown in the figure below.



When you choose the **Pie** chart, a plot will be created for every sample selected in the active sample list. Before using this option, ensure that only the samples you want to plot are selected.

You must create a generic pie plot for every sample before you can customize it for each individual one. To do so, in the initial options dialogue set up the general options for the **Pie** plot - parameters to be plotted, their color and pattern, etc. Once this is done, click **[OK]**. A pie plot will be created for every selected sample. Right click on any of the pie plots to launch the **Pie Plot Options** dialogue and use it to customize the plots for each sample.

Sample





At the top of the **Pie Plot Options** dialogue, there is a **Sample** field which identifies the active sample(s) associated with the plot options dialogue. If you have created several Pie charts, you can use the **Sample** field to select and modify the settings for each pie plot individually, or several at once as shown below.

Sample				
ID	SAMPLEID	LOCATION	WATERTYPE	SAMPLE_DAT
1	MW-1-92	Waterloo	Ca-Na-SO4-Cl	8/15/199
2	MW-1-93	Waterloo	Ca-Na-SO4-Cl-HCO3	6/1/199
3	MW-1-94	Waterloo	Ca-Na-SO4-Cl	6/15/199

To make changes to multiple plots simultaneously, simply select all of the samples you wish to edit in the list above (using your mouse), make the required plot changes then click **[Apply]**. This is useful for normalizing the axes scales, units, and general formats for each of the open Pie charts. If just one sample is highlighted and selected in this dialogue, then the changes will only be applied to this plotted sample.

Parameters frame

Parameters list

The **Parameters** list contains the parameters that will be plotted on the Pie chart. Existing parameters can be changed by selecting the parameter, and then typing in the name of the new parameter in the same field. New parameters can be added to the list by clicking the  button and selecting a parameter from the available list. The new parameter will then be added to the bottom of the list. Parameters can be removed from the list by selecting the parameter and clicking the  button. The order of the parameters in the list can be adjusted using the  and  buttons.

Fill Style frame

The **Fill Style** allows you to edit the appearance of each pie slice in the plot. Select a parameter from the Parameters list, then select the **Pattern** and the **Color** you wish to assign to that slice of the pie chart. Repeat for the other parameters in the list.

Unit frame

Allows you to set display unit. The typical unit is meq/l which places cations in the upper part of the plot and anions in the lower part or vice versa, and permits a visual control of the electroneutrality.


Inside Circle frame

The **Inside Circle** is used to represent the concentration of a single parameter (or combination of parameters) that is not included in the **Pie** chart parameters. You can specify any chemical or physical parameter that reveals the most comprehensive information about the sample. For example, you may want to display the CO₂ or SiO₂ concentration for the selected sample. Type a valid parameter in the **Proportional to** field to create an inside circle on your pie chart. Type a **Maximum Value** in the next field below (or use the default value). If several pie charts are created, it is recommended that you choose 3 to 4 times the average value of the inside parameter as the maximum value. All values that are greater than or equal to the maximum value are displayed with the maximum diameter. The inside circle will have a radius between zero and the specified maximum radius (R).

- R = Maximum radius for the internal circle
- Value = value for this sample for the selected parameter.
- Maximum Value: Maximum value above which, the full internal circle is drawn.
- Inside Radius = $(\text{Value} / \text{Maximum Value}) * (\pi R/2)$

Unless the parameter you have selected is unitless, you must also specify the units using the **Units** combo box.

Plot frame

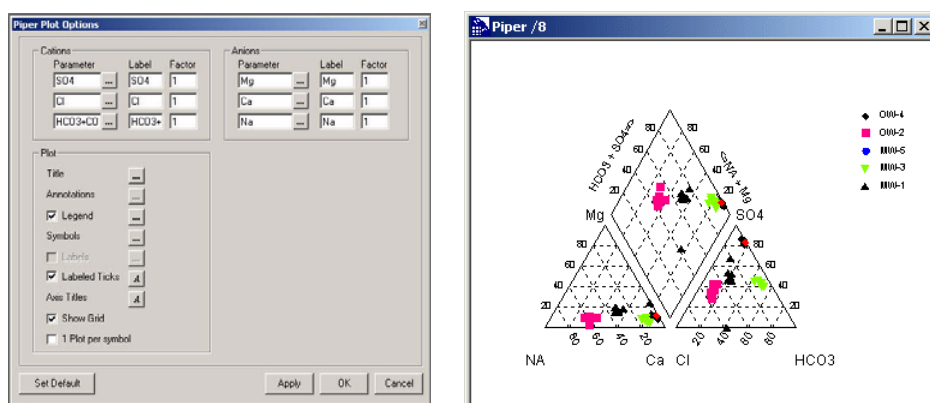
Click the  button beside **Title** and **Legend** to load the corresponding options dialogues. Pie plots are titled automatically according to the settings in **File>Preferences>Plots**. The **Legend** feature must be activated (checked) before the settings dialogue can be accessed.

Slice labels frame

The **Slice Label** options will apply a label to each slice of the pie chart. You can choose from **Parameter Name**, **Concentration**, **Percentages**, or **None** for no label.

4.5.12 Piper Plot

The **Piper** plot is useful for showing multiple samples and trends in major ions. An example of the **Piper** plot and the corresponding **Piper Plot Options** dialogue is shown in the figure below.



In the **Piper** plot, major ions are plotted as cation and anion percentages of milliequivalents in two base triangles. The total cations in meq/l, and the total anions in


meq/l, are set equal to 100%. The data points in the two triangles are then projected onto the diamond grid. The projection reveals certain useful properties of the total ion relationships. Every sample is represented by three data points; one in each triangle and one in the projection diamond grid.

If your database stores carbonate species concentration as alkalinity, you may use the parameters for alkalinity instead of HCO_3 . To do this add CO_3 to the field so that it displays HCO_3+CO_3 .

The **Piper** plot allows comparisons of 6 parameters between a large number of samples. Like all trilinear plots, it does not portray absolute ion concentrations. The main purpose of Piper plots is to show clustering of samples.


Cations and Anions frames

The default parameter settings include the major cations (Na, Ca, and Mg) against the major anions (Cl, SO_4 and HCO_3); however, any other parameters, or combination of parameters, can be easily selected (e.g. gas composition, trace elements etc.)

The parameters can be modified by typing the Internal name of the parameters in the appropriate fields, or pressing the  button and selecting another parameter from the available list. You may also multiply or divide the parameter concentrations by a factor if you need to compare elements which are systematically very different in concentration (e.g. comparing a trace element together with major ions). Finally, you may also enter parameter operators and functions (e.g. $\text{Na}+\text{K}$ or Na/Cl); to do this, simply enter $\text{Na}+\text{K}$ (for example) in the parameter field manually, and press **[Apply]**. However multiple parameters (e.g. Na, Cl) are not accepted.

Both the Cations and Anions frames also contain a **Label** field for each parameter. This allows customization of the parameter labels on the plot diagram. For example the parameter 'Ca' can be edited in the **Label** field to displayed as 'Calcium'.

Plot frame

Use the  link buttons in this frame to modify the options for the **Title**, **Legend**, **Symbol**, **Labels**, and **Format**.

Click the 1 Plot per Symbol option to create one Piper plot for each station.

4.5.13 Probability plot


Probability plots involve plotting quantiles of a data against the quantiles of the normal distribution. They are particularly useful for spotting irregularities within the data when compared to a specific distributional model, like the Normal. It is easy to determine whether departures from Normality are occurring more or less in the middle ranges of the data or in the extreme tails. **Probability** plots can also indicate the presence of possible outlier values or a different population that do not follow the basic pattern of

the data and can show the presence of significant positive or negative skewness. The log scale transformation on the y-axis may be used to test whether a dataset that shows non normal behavior is simply normally distributed.

X-Axis frame


Use the  button beside **Format** to set the Y-axis options.

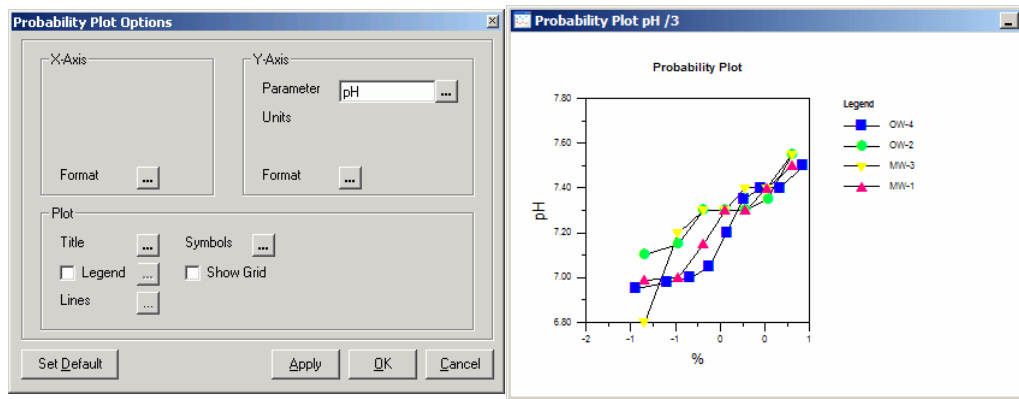
Y-Axis frame

Select the parameter to plot by clicking the  button and choosing from the parameter list. If the chosen parameter has units, a **Units** combo box will be presented in which you can select the desired units.

Use the  button beside **Format** to set the X-axis options.

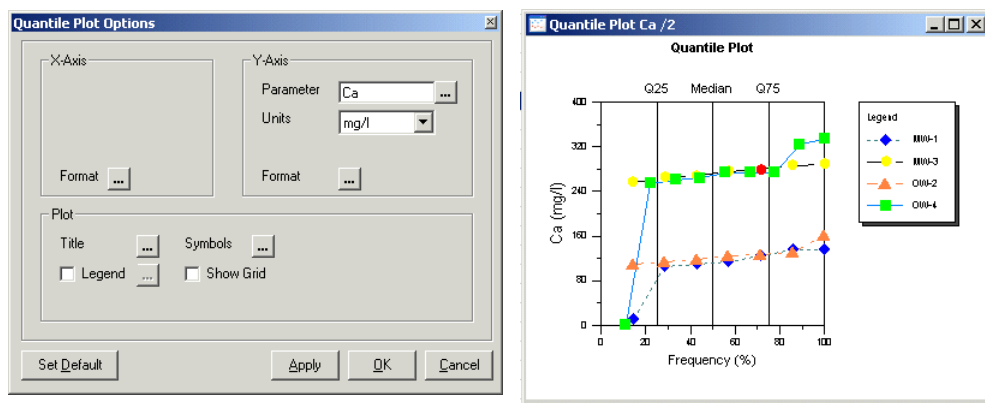
Plot frame

Use the  button beside **Title**, **Legend**, **Symbol** and **Lines** to set their options. Check the box beside **Show Grid** to display the gridlines on the plot. Creating correlation lines may be useful to visually test whether data is distributed on a straight line, indicating a normal distribution of the underlying data set. if the y-axis is set to log scale, then the alignment of the data on a line indicates lognormal distribution.



4.5.14 Quantile Plot


The **Quantile** chart plots the ranked data from lowest to highest on the background of lines that denote 25% Quartile, Median, and 75% Quartile.



X-Axis frame


Use the  button beside **Format** to set the Y-axis options.

Y-Axis frame

Select the parameter to plot by clicking the  button and choosing from the parameter list. If the chosen parameter has units, a **Units** combo box will be presented in which you can select the desired units.

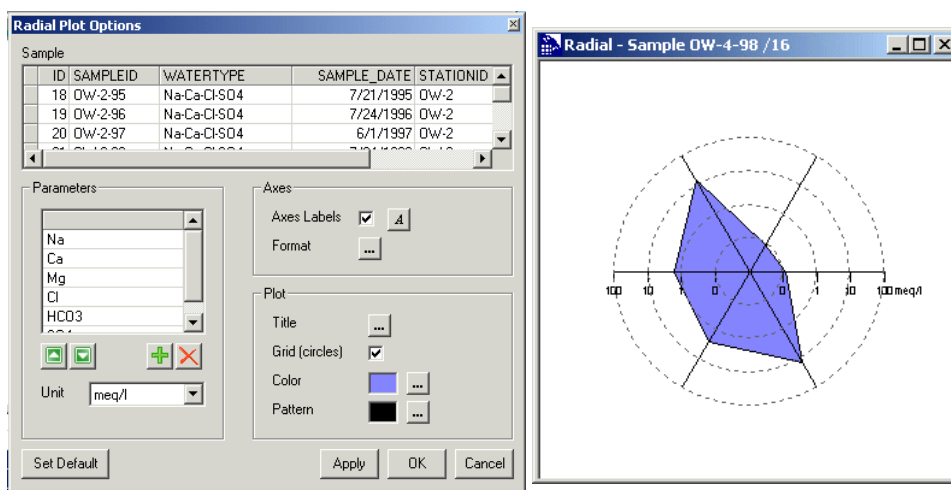
Use the  button beside **Format** to set the X-axis options.

Plot frame

Use the  button beside **Title**, **Legend**, and **Symbol** to set their options. Check the box beside **Show Grid** to display the gridlines on the plot.

4.5.15 Radial Plot

The **Radial** plot is used to compare multiple parameter values for a single sample and to compare the ratios of these values for many different samples. An example of the **Radial** plot and the corresponding **Radial Plot Options** dialogue is shown in the figure below.



Radial plots can be used to evaluate the change in water quality at a single location over a period of time, or they can be used to evaluate the change in water quality as the water passes through different geologic formations or different subsurface conditions. A plot will be created for every selected sample in the Active Samples List. Before using this option, ensure that only the samples you want to plot are selected.




Sample

The **Sample** field identifies the active sample(s) associated with the plot options dialogue. If you have several **Radial** plots open, you can use the **Sample** field to select and modify the settings for each one.

To make changes to multiple plots simultaneously, simply select all of the samples in the list above (using your mouse), make the required plot changes, then click **[Apply]**. This is useful for normalizing the axes scales, units, and general formats for each of the open **Radial** plots. The changes apply to most of the graph settings that affect the appearance of the **Radial** plot, with the exception of the plot title. If just one sample is highlighted and selected in this dialogue, then the plot options changes will only be applied to this plotted sample.

Parameters frame

The **Parameters** list contains the parameters that will be plotted on the **Radial** plot. A **Radial** plot requires a minimum of three parameters. Existing parameters can be changed by selecting the parameter, and then typing in the name of the new parameter in the same field. New parameters can be added to the list by clicking the **+** button and selecting a parameter from the available list. The new parameter will then be added

to the bottom of the list. Parameters can be removed from the list by selecting the parameter and clicking the  button. The order of the parameters in the list can be modified by selecting the parameter and using the  or  buttons to move the position of the selected parameter.


Use the **Units** combo box to choose the appropriate units.

Axes frame

Checking the **Axes labels** box will display the respective parameter labels at the outer ends of each axis.

Click the  button beside **Format** to set the axes ticks options.

Filling options frame

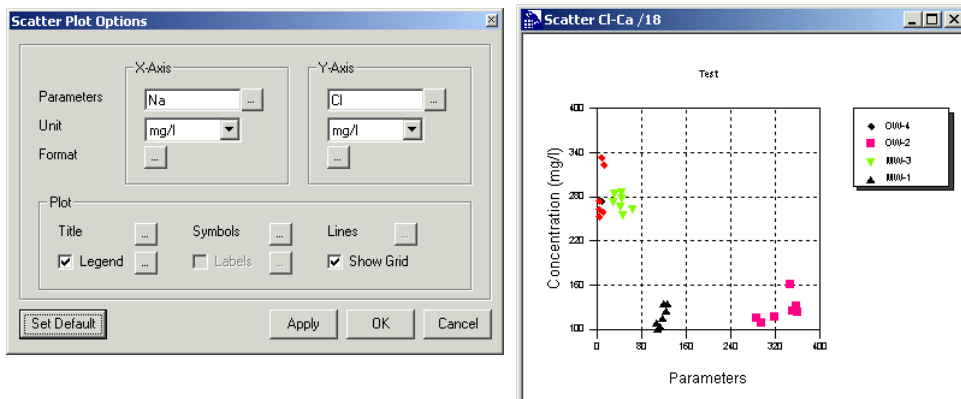
Click the  button beside **Title** to create a title for the plot. Whereas many of the options are modified in all plots when applied, the title will be created unique for the active plot, unless you have several plots selected.

Use the **Grid (Circles)** check box to show/hide the circular grid of the plot.

Use the  button beside **Color** and **Pattern** to set these respective features.

4.5.16 Scatter Plot

X-Y Scatter plots are the most simple and popular approach to interpreting hydrochemical data. **Scatter** plot shows effects such as correlation of parameters or clustering of samples in a very intuitive manner. An example of the **Scatter** plot and the corresponding **Scatter Plot Options** dialogue is shown in the figure below.



X-Axis and Y-Axis frames

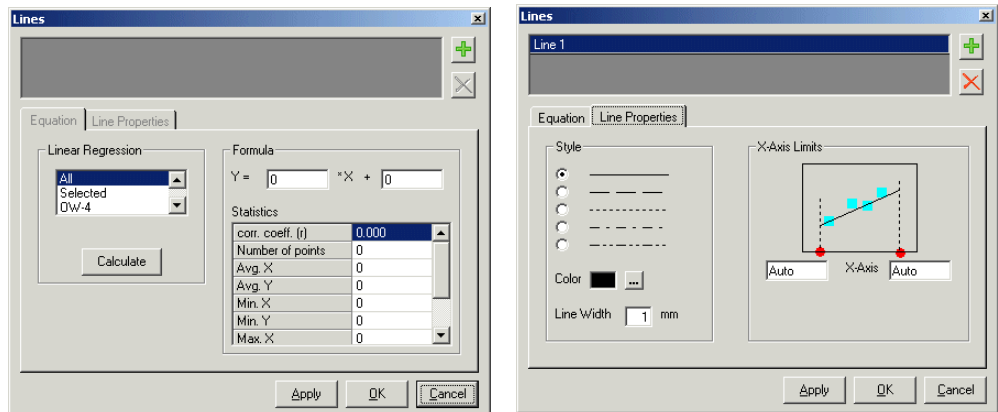
Click the button beside **Parameter** field to select the parameter to plot on the respective axis. Unless the chosen parameter is unitless, the **Unit** field will be enabled. Use this combo box to select the appropriate units for the chosen parameter.

Once you have defined the parameter options, click on the button beside **Format** to set the axis options.

Plot frame

Click the button beside **Title**, **Legend**, **Symbol**, **Label**, and **Lines** to customize their respective settings.

The **Scatter** plot is one of several plots in AquaChem for which you can add custom trend lines. To do so, click on the button besides **Lines**. Use the dialogues shown below to create and customize the trend line:



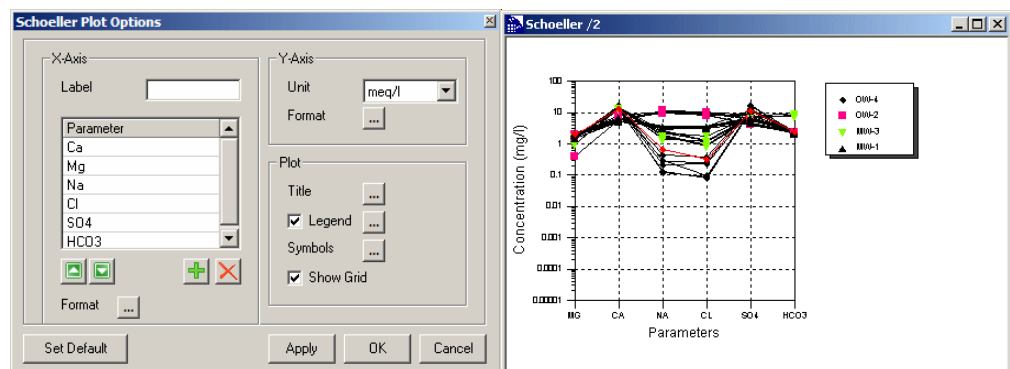
The **Lines** options allow you to perform a regression analysis on all the data, or on a specified group of data. You can calculate the equation of best fit for the line, and plot the line on the graph. You can also create your own line with your own equation and plot it on the graph as well.

For more information on how to create a line on the plot, please refer to “Line Dialogue” on page 185.





This is one of the plots on which it is common to use precision bars. You can do so using the Symbols Dialogue described on page 182 of this chapter.

4.5.17 Schoeller Plot

Schoeller (1962) developed semi-logarithmic plots to represent major ion analyses in milliequivalents per liter and to demonstrate different hydrochemical water types on the same plot. The number of analyses that can be illustrated at one time is limited because of the lines. The plot has the advantage that, unlike trilinear plots, actual parameter concentrations are displayed. An example of the **Schoeller** plot and the corresponding **Schoeller Plot Options** is shown in the figure below.



X-Axis frame

Customize the list of parameters to be plotted. New parameters can be added to the list by clicking the  button and selecting a parameter from the list of available parameters. The new parameter will then be added to the bottom of the list. Parameters can be removed from the list by selecting the parameter and clicking the  button. The order of the parameters in the list can be modified by selecting the parameter and using the  or  buttons to move the position of the selected parameter. Parameters can include ratios, sums, or differences (Na/Cl, Ca+Mg, Cl-SO4). Simply manually enter these operators in the parameter field.


Click the  button beside **Format** to set the axis options.

Y-Axis frame

The y-axis for the **Schoeller** plot is concentration of the chosen parameters. The **Unit** combo-box will allow you to choose the appropriate units. If ions are portrayed in a **Schoeller** plot, then it is recommended to use the meq/l unit, since it allows to compare directly the ion ratios.

Click the  button beside **Format** to set the axis options.

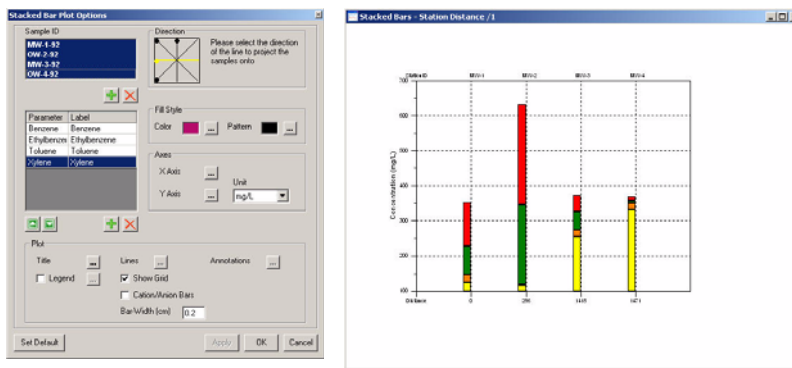
Plot frame

Click the  button beside **Title**, **Legend**, and **Symbols** to customize their respective settings.

Check the **Show Grid** box to display the gridlines on the plot.

4.5.18 Stacked Bars (Distance)

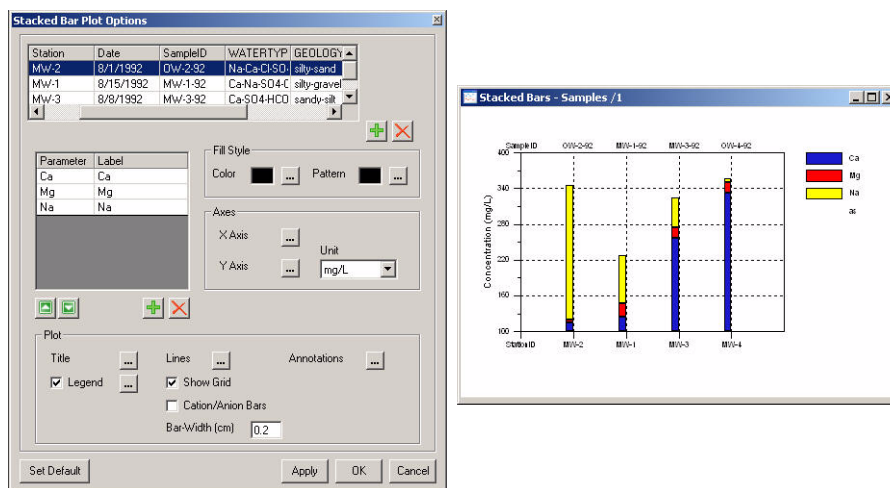
This plot option allows you to display stacked bars for multiple stations which are projected onto a line. Generally this is intended for only one sample per station. The x axis will display the distance between the stations to allow users you make spatial interpretations.



In the Options dialogue box you will find the list of samples you selected to be included in your plot. You can add or remove samples from this list with the green plus or red x respectively. You need to indicate the direction of the line that the samples will be projected onto.

4.5.19 Stacked Bars (Stations)

This plot option allows you to select which stations you wish to plot stacked bars for. Generally this is intended for displaying one sample per station. Where the time series (Stacked Bars) option will display all samples for the selected station, this plot allows you to pick which sample you wish to plot for each station. This allows you to make cross station comparisons.

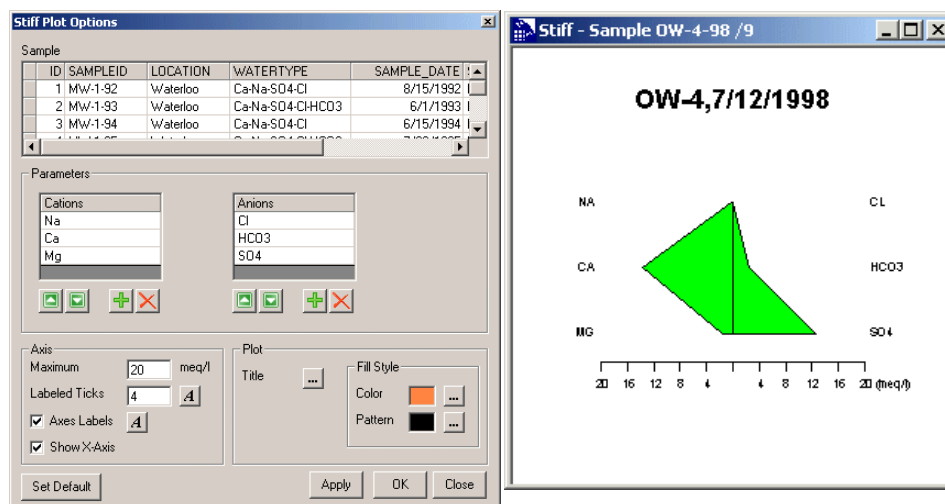


In the Options dialogue box you will find the list of samples you selected to be included in your plot. You can add or remove samples from this list with the green plus or red x respectively.

4.5.20 Stiff Plot

The **Stiff** plot belongs to the group of pattern plots (see Hem 1985, p. 175). It is constructed by plotting the milliequivalents per liter of three or more anions and three or more cations. Stiff plots can be used to evaluate the change in water quality at a single location over a period of time, or they can be used to evaluate the change in water quality as the water passes through different geologic formations or different subsurface conditions.

An example of the **Stiff** plot and the corresponding **Stiff Plot Options** dialogue is shown in the figure below.



A **Stiff** plot will be created for every selected sample in the active list. Before using this option, ensure that only the samples you want to plot are selected. The following section describes some of the features and options of the **Stiff** plot that are not covered in the Common Plot Features section.





Sample

The **Sample** field identifies the active sample(s) associated with the plot options dialogue. If you have several **Stiff** plots open, you can use the **Sample** field to select and modify the settings for each one.

To make changes to multiple plots simultaneously, simply select all of the samples in the list above, (using your mouse), make the required plot changes, then click **[Apply]**. This is useful for normalizing the axes scales, units, and general formats in each of the open **Stiff** plots.

Parameters frame

The **Cations** and **Anions** lists contain the parameters that will be plotted on the **Stiff** plot. Existing parameters can be changed by selecting the parameter, and then typing in the name of the new parameter in the same field.

New parameters can be added to the list by clicking the  button and selecting a parameter from the list of available parameters. The new parameter will then be added to the bottom of the list. Parameters can be removed from the list by selecting the parameter and clicking the  button. The order of the parameters in the list can be modified by selecting the parameter and using the  or  buttons to change the position of the selected parameter.


NOTE: The **Stiff** plot requires an equal number of cations and anions (i.e. typically three cations and three anions).

Axis frame

Set the **Maximum** concentration for the x-axis in meq/l.

Set the number of **Ticks** for the x-axis.

Use the **Axes Labels** and **Show X-Axis** check boxes to show/hide the respective features.

The font for the tick and axis labels can be changed by clicking the  button.

Plot frame

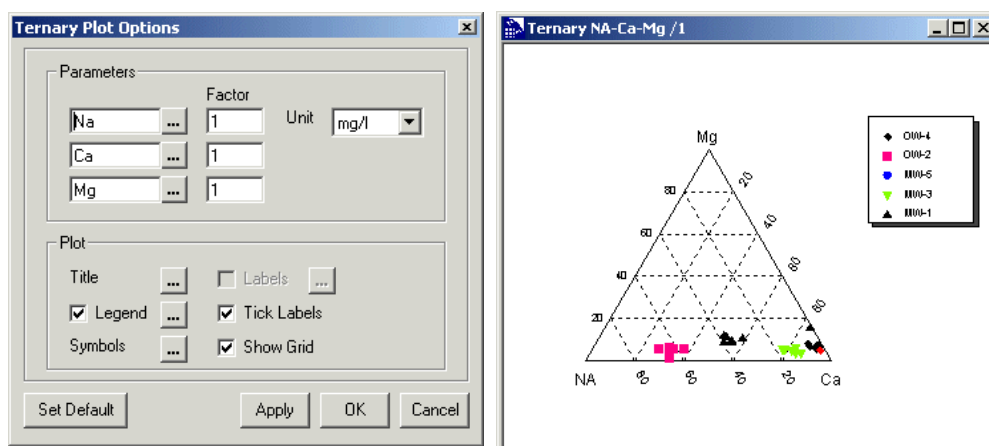
Use the  button beside **Title** and **Legend** to customize their settings.

Fill Style frame

Access the fill **Pattern** and the **Color** options by clicking the  button.


4.5.21 Ternary Plot

Ternary plot is used to determine the relationship between the concentrations of three different parameters in multiple samples. An example of the **Ternary** plot and the corresponding **Ternary Plot Options** dialogue is shown in the figure below.



Like the **Piper** and **Durov** plots, the **Ternary** plot displays relative concentrations of each parameter with respect to the sum of the concentrations of each parameter. Each vertex of the Ternary plot represents a relative concentration of 100% for the parameter at the respective vertex, while the base represents a relative concentration of 0% for the parameter plotted at the opposite vertex.


Parameters frame

The parameters can be modified by typing the Internal name of the parameters in the appropriate fields, or pressing the  button, and selecting another parameter from the dialogue that appears.

You may also multiply the parameter concentrations by a factor if you need to compare elements which are systematically very different in concentration (e.g. comparing a trace element together with major ions). You may also enter parameter operators and functions (e.g. Na+K or Na/Cl); to do this, simply enter Na+K in the parameter field and press **[Apply]**. However multiple parameters (e.g. Na,Cl) are not accepted.

The **Unit** combo box will allow you to select the appropriate units (where applicable).

Plot frame

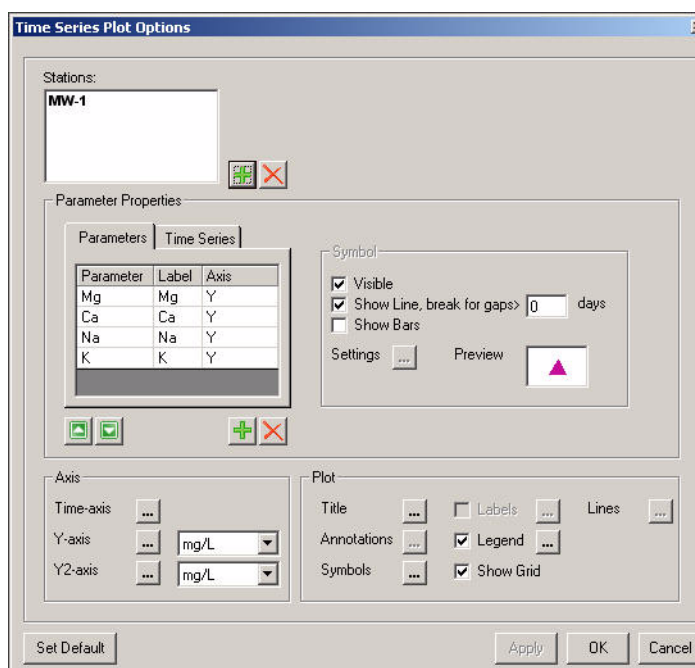
Click the  button beside **Title**, **Legend**, **Symbol**, and **Labels** to customize their respective settings. Use the **Show Grid** check box to show/hide the plot gridlines.

4.5.22 Time Series (Multiple Parameters)

The **Time Series** plot shows the evolution of a chemical or physical parameter for a given sampling point as a function of time. This plot is a standard technique for interpreting hydrochemical and hydrogeological processes and in particular temporal trends in natural waters.

Time Series (Multiple Parameters) plot is intended for one sampling station (or any other defined legend item, e.g. aquifer).

The options dialogue for the Time Series (Multiple Parameters) plot appears below.



Choose a single **Station** (or other defined legend item) from the top of the dialogue box.

Parameter Properties frame


Parameters Tab

Choose the parameter(s) that you want plotted. Use the  and  buttons to add or remove parameters, and the  and  buttons to change the order of the parameters.

The **Name** of the parameter is automatically filled in, however you may manually change it under the **Label** column. Select the desired Y-axis under the **Axis** column (Y refers to the left Y axis, and Y2 refers to the right Y axis)

Time Series Tab


The time series tab allows you to select non-chemical time series data to show on the time series plot, e.g., precipitation, water levels etc. Please note that you must import the time series data into AquaChem database before you can show on the time series plot. For more information on importing time series data, please refer to “Import” on page 52.

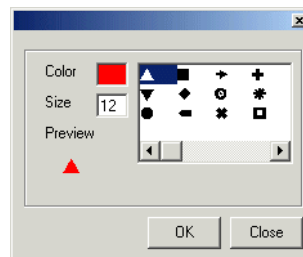
When the Time Series Tab is selected, click the  button to load a time series dataset. The Parameters window will appear on your screen. From the Category combo box select the table in which the time series is saved. The available time series data series for the selected table will be displayed. Choose the desired time series and click the **[Select]** button.

The **Name** of the parameter is automatically filled in, however you may manually change it under the **Label** column. Select the desired Y-axis under the **Axis** column (Y refers to the left Y axis, and Y2 refers to the right Y axis)


Symbol frame

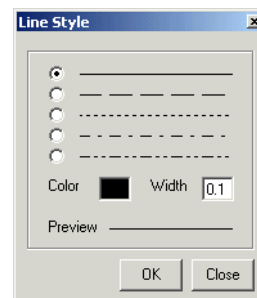
Use the **Visible** check box to show/hide symbols.

Below the **Visible** check box is a preview of the symbol used to represent currently selected parameter on the plot. Click the  button beside the preview to load the dialogue shown to the right. Use this dialogue to select the symbol style, size, and color (click on the field beside **Color** to produce the color selection dialogue). Click **[OK]** when satisfied with symbol settings.




Line frame


Use the **Visible** check box to show/hide the line connecting the symbols. Click the  button beside the line preview to produce the dialogue shown on the right. Use this dialogue to set the style, color, and width for the line (click on the field beside **Color** to load the color selection dialogue). When you are satisfied with the line settings, click **[OK]**.



Axis frame

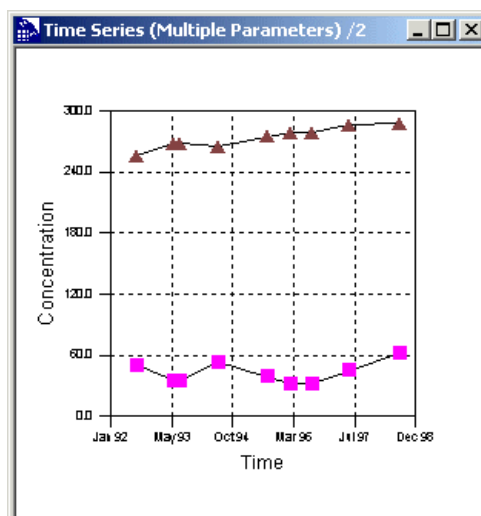
Click the  button beside **Time-axis** and **Y-axis** to customize their respective settings.

Plot Frame

Click the  button beside **Title**, **Legend**, **Symbol**, and **Label** to customize their respective settings. The **Legend** and **Label** features must be activated (checked) before their dialogues can be accessed. Use the **Show Grid** check box to show gridlines on the plot.

This is another plot where precision bars may be useful. For more information on how to add them, please refer to “Symbols Dialogue” on page 182.

The figure below demonstrates a finished **Time Series (Multiple Parameter)** plot:

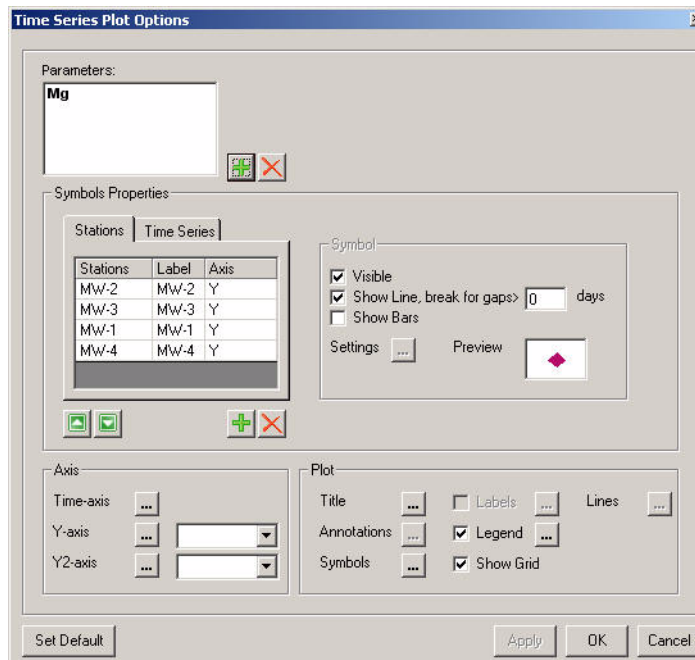


4.5.23 Time Series (Multiple Stations)

The **Time Series** plot shows the evolution of a chemical or physical parameter for a given sampling point as a function of time. This plot is a standard technique for interpreting hydrochemical and hydrogeological processes and in particular temporal trends in natural waters.





Time Series (Multiple Stations) is intended for one parameter.

The options dialogue for the Time Series (Multiple Stations) plot appears below.



Choose a single **Parameter** from the top of the dialogue box.

Station Properties frame

Choose the station(s) that you want plotted. Use the  and  buttons to add or remove stations, and the  and  buttons to change their order.


Use the **Unit** combo box to select the appropriate units.

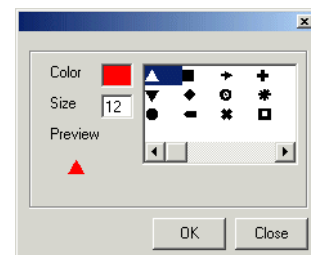
The **Name** of the station is automatically filled in, however you may manually change it.

Symbol frame

Use the **Visible** check box to show/hide symbols.


Below the **Visible** check box is a preview of the symbol used to represent currently selected station on the plot.

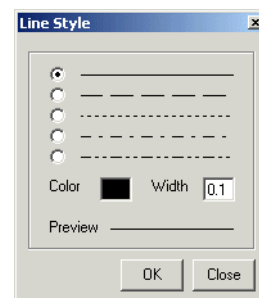
Click the  button beside the preview to load the dialogue shown to the right. Use this dialogue to select the symbol style, size, and color (click on the field beside **Color** to produce the color selection dialogue). Click **[OK]** when satisfied with symbol settings.




Line frame

Use the **Visible** check box to show/hide the line connecting the symbols.


Click the  button beside the line preview to produce the dialogue shown on the right. Use this dialogue to set the style, color, and width for the line (click on the field beside **Color** to load the color selection dialogue). When you are satisfied with the line settings, click **[OK]**.



Axis frame

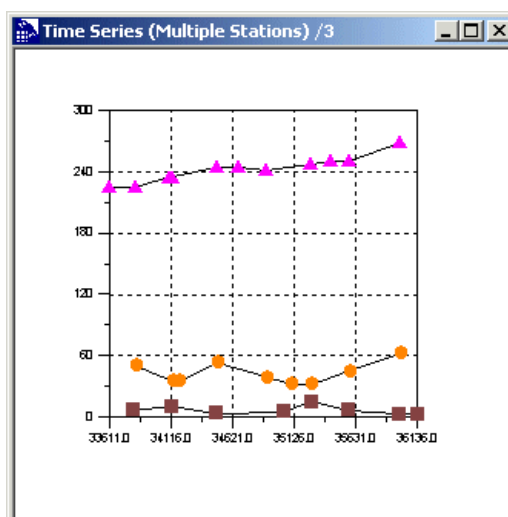
Click the  button beside **Time-axis** and **Y-axis** to customize their respective settings.

Plot Frame

Click the  button beside **Title**, **Legend**, **Symbol**, and **Label** to customize their respective settings. The **Legend** and **Label** features must be activated (checked) before their dialogues can be accessed. Use the **Show Grid** check box to show gridlines on the plot.

Please refer to “Symbols Dialogue” on page 182 for information on how to add precision bars.

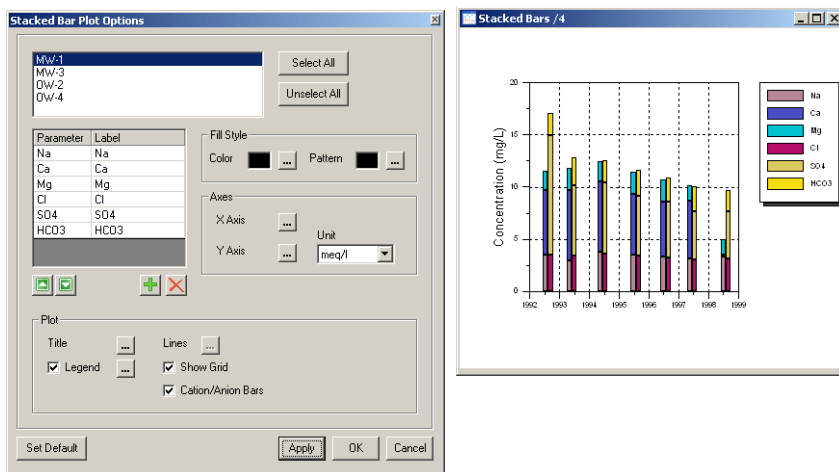
A finished **Time Series (Multiple Stations)** plot is shown below:





4.5.24 Time Series (Stacked Bars)

The stacked bar chart allows you to display the chemical concentrations of samples as stacked bars. Major anion and cation bars can be stacked side-by-side to show the ion balance over time. Different heights for anion and cation bars can indicate an ion imbalance, and point out an issue in a particular sample.

An example of the **Stacked Bar** plot and the corresponding chart options are shown in the figure below.



Parameter Frame

By default, the major ions are listed in the parameter frame. However, you may add and remove parameters as desired using the  and  buttons. The Label column in the parameter list allows you to change the labels of each parameter for displaying on the plot legend, e.g., Na to Sodium.

Fill Style Frame

Specify the fill color and pattern of the stacked bar for the selected parameter.

Axis Frame

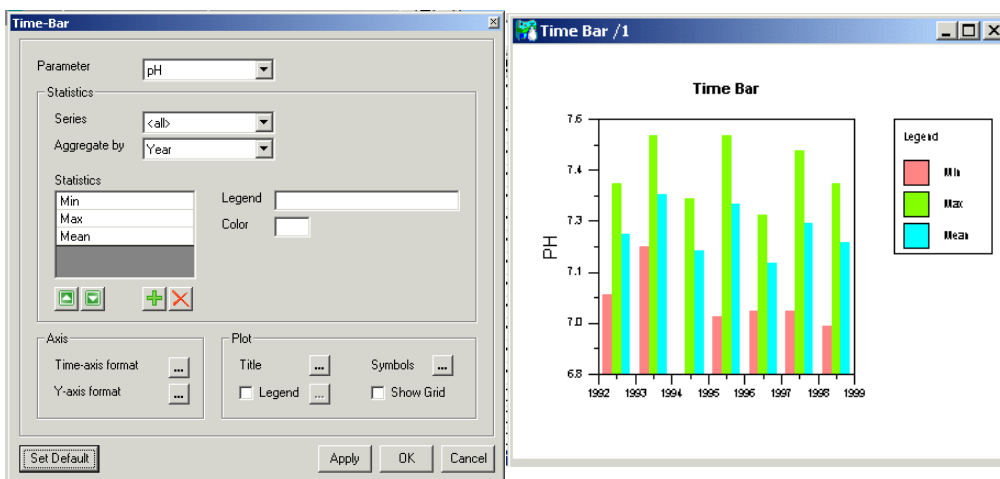
Click the [...] button to customize the X-axis and Y-axis settings. Use the **Unit** combo box to select the appropriate units.

Plot Frame

Click the [...] button beside **Title** and **Legend** to customize their respective settings. Use the **Show Grid** check box to show gridlines on the plot. Use the **Cation/Anion Bars** check box to show the cations and anions in two separate bars.

4.5.25 Time Series (Statistics Summary)

Time Series (Statistics Summary) plot is used to display a statistical summary over a specified time period as a bar chart. The options for the time period over which the data is to be aggregated include month or year. The following summary values per period can be displayed: as **Min**, **Max**, any **Quantile** and **Standard Deviation**.



Parameter



Select the parameter for which you wish to create the **Time Bar** plot, and specify the appropriate units



Statistics frame

Select the **Series**, the values from which will be plotted.

Select the time period by which to **Aggregate** the data.

Select the **Statistics** to display on the plot. Minimum, Maximum, and Mean are selected by default. To change a statistic, click on the statistic you wish to change and

choose a new one from the presented combo box. To add a new statistic, click the  button; to delete an existing statistic, click the  button.

Use the  and  buttons to change the order the statistic bars appear on the plot.


Legend field shows the statistic label that will be displayed in the plot legend for the statistic currently selected. If you wish a different label to appear for this statistic, simply replace the default text with new text.

Color field shows the color of the bar for the statistic currently selected in the **Statistics** list. To change the color, double-click on the field to load the color selection dialogue.

Axis frame

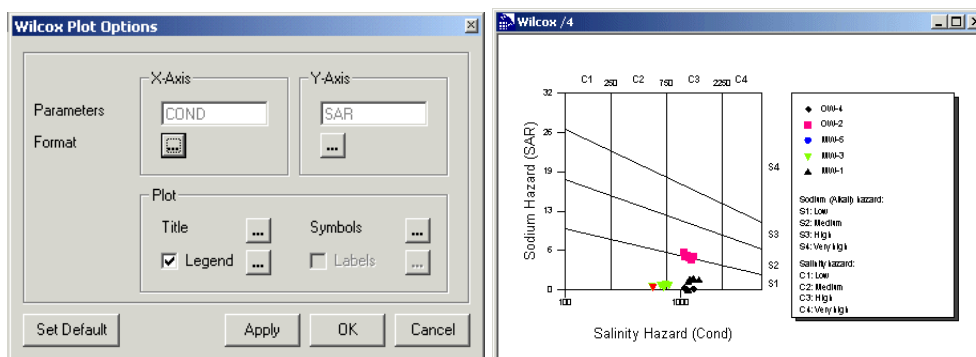
Click the  button beside **Time-axis format** and **Y-axis format** to customize their respective settings.

Plot Frame

Click the  button beside **Title**, **Legend**, and **Symbol** to customize their respective settings. Use the **Show Grid** check box to show gridlines on the plot.


4.5.26 Wilcox Plot

A **Wilcox** plot can be used to quickly determine the viability of water for irrigation purposes. The **Wilcox** plot is also known as the U.S. Department of Agriculture diagram. An example of the **Wilcox** plot and the corresponding **Wilcox Plot Options** dialogue is shown in the figure below.




The **Wilcox** plot is a simple scatter plot of **Sodium Hazard (SAR)** on the **Y-axis** vs. **Salinity Hazard (Cond)** on the **X-axis**. The Conductivity (**COND**) is plotted by default in a log scale. These plot parameters are hardwired into AquaChem and cannot be changed. You must have values for the COND and SAR parameters in your database in order to use this plot.

X-Axis and Y-Axis frames

The Parameters are pre-selected and cannot be changed. Click the  button to customize the x-axis and y-axis settings.

Plot frame

Click the  button beside **Title**, **Legend**, **Symbols**, and **Labels** to customize their respective settings. The **Legend** and **Labels** features must be activated (checked) before their options dialogues can be accessed.

The **Wilcox** plot contains the following sections:

Conductivity (us/cm)

C1: Low (0-249)

C2: Medium (250-749)

C3: High (750-2249)

C4: Very High (2250-5000)

The SAR values are divided into the following categories:

S1: Low

S2: Medium

S3: High

S4: Very High

The locations of the SAR lines are determined by the following equations:

S1: Line equation: $y = -1.5816e-3x + 10.15816$

S2: Line equation: $y = -2.2959e-3x + 18.22959$

S3: Line equation: $y = -3.0102e-3x + 26.30102$

5

Reports

When you select **Reports** from the main menu, you are presented with several types of reports to aid in the presentation of your data. These reports are generated in a separate **Report** window as unformatted text. These reports can be printed ‘as is’ using the **[Print]** button located on the lower-left corner of the report window, or the information can be saved using the **[Save]** button.

The following pre-defined Reports are included with AquaChem:

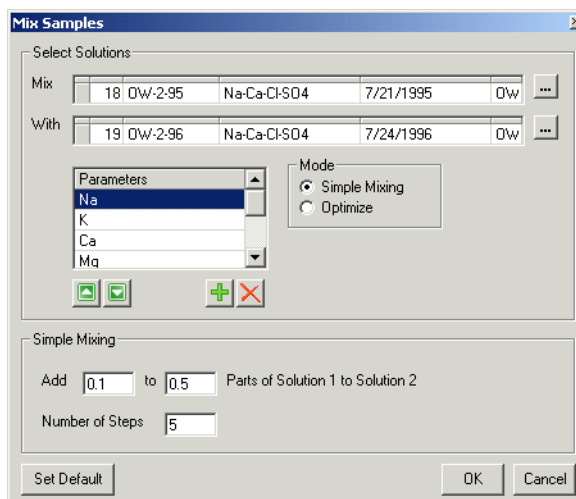
- Data Summary
- Compare Sample
- Mix Samples
- Water Quality Standards
- Hardness Dependent Standards
- Rock Source Deduction
- Statistics
 - Summary Statistics
 - Correlation Matrix
 - Trend Analysis
 - Outlier tests
 - Tests for Normality
 - Alert Levels

You can also design your own report templates, which allows you to produce reports on a wide variety of sample data and calculations. You can do so using the **Report Designer** component, which is explained later on in this chapter. AquaChem includes a **Sample Summary Report**, which was designed using the Report Designer.

5.1 Common Report Features

Although each pre-defined report has unique characteristics, there are also a few buttons and options that are common for all reports. These buttons may be found below the **Parameters** list in the **Report Options** dialogue, or in the **Report** window itself.

A generic Report options dialogue is shown below:



The **Parameters** list is common to many reports dialogues. Below the **Parameters** list you will find the following buttons:



The up arrow button allows you to move the selected parameter up in the report options dialogue.



The down arrow button allows you to move the selected parameter down in the report options dialogue.




The sort button allows you to sort the parameter list alphabetically.

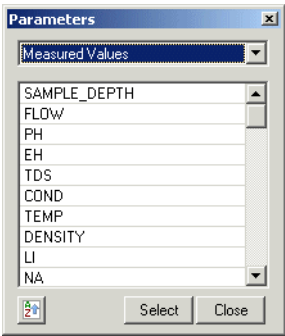


The delete button will remove the selected Parameter from the parameters list.



The add button allows you to select a parameter from the list of available parameters and add new parameters to the Report. When you press the add button, the following dialogue will appear:

Simply choose the desired parameter and press the **[Select]** button, and this parameter will be included in the report. You can use the  button to sort the parameter list alphabetically, allowing you to quickly locate a parameter. In addition, some reports allow you to choose from several parameter categories, which are available in the combo box at the top of the **Parameters** dialogue. Once you are finished, press the **[Close]** button to close this dialogue.

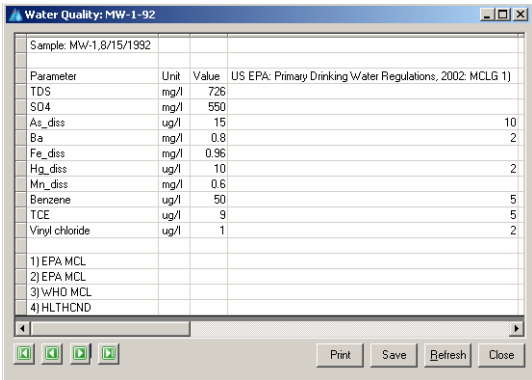


When the **[Set Default]** button is pressed, the current parameter settings will be used as the default parameter settings for all new reports for the selected report type.



The **Select sample** button loads your list of active samples, and allows you to select a sample for the Report. **Double-click** on the desired sample to load this into the Report options.

A generic report window is shown below:



In some **Report** windows, you will find the following buttons:



The scroll buttons allow you to generate a Report for other samples in your active list. These buttons can be found on the bottom of the Report window. The function of these buttons (in order from left to right) is as follows:

First sample - loads a Report for the first sample in your active list.

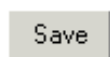
Previous sample - loads a Report for the previous sample in your active list.

Next sample - loads a Report for the next sample in your active list.

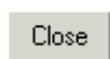
Last sample - loads a Report for the last sample in your active list.



The **[Print]** button will print the current report to the selected Windows printer.



The **[Save]** button will save the current report. The pre-defined reports in AquaChem can be saved as .TXT, .CSV, and .XLS. The customized reports (designed using the **Report Designer**) can be saved as .HTM or .RTF format.



The **[Close]** button will close the Report window.

Where applicable, you can access the Report options dialogue by selecting **View** from the main menu and then **Options**. This allows you to change the report settings (such as which parameters or samples to use).

The following section provides more details on each of the various pre-defined Reports available in AquaChem.

5.2 Data Summary

The **Data Summary** report allows you to generate a summary report for the AquaChem database and all stations currently displayed in the Active Samples list. The following information is shown on a data summary report:

- Database location path
- Database Template
- Active Stations
- Active Samples
- Number of Parameters
- Number of samples per station including the first sample and last sample dates.
- All analyzed parameters in the samples, including the units, results, %ND, MDL, Min and Max.

- Results per Station matrix showing the number of measurements for each parameter and station combination.

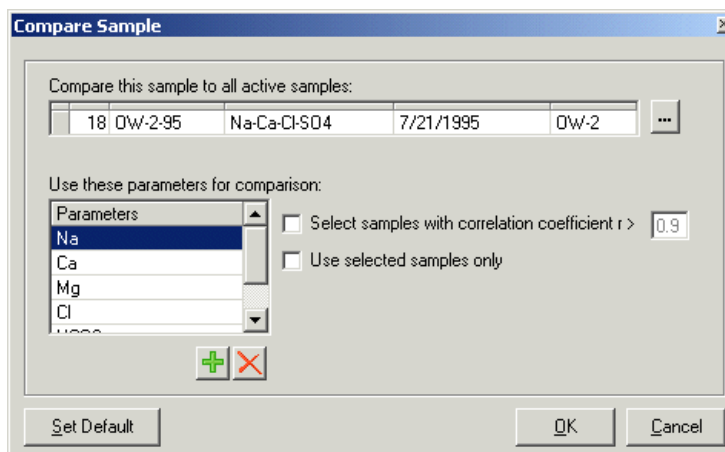
Database Summary						
Database	D:\DOCUMENTS\AquaChem\Demo.aqc					
Database Template	0					
Active Stations	4					
Active Samples	30					
Parameters	82					
Stations						
Name	Samples	First sample	Last sample			
MW-1	7	8/15/1992	8/1/1998			
MW-2	7	8/1/1992	7/24/1998			
MW-3	7	8/8/1992	7/30/1998			
MW-4	9	7/15/1992	7/12/1998			
Parameters						
Name	Unit	Results	%ND	MDL	Min	Max
pH_field		30	0.00		0	15.7
K	mg/l	30	0.00		6.8	7.55
Na	mg/l	30	1.69	1	1	2.5
Ca	mg/l	30	1.69	2	2	269
Mg	mg/l	30	1.69	10	1.6	333
Fe_diss	mg/l	30	2.17	1	1	26.9
Cl	mg/l	17	2.17	0.15	0.02	0.96
HCO3	mg/l	30	1.67	1	1	361
Li	mg/l	30	0.00		1.2	601
Si	mg/l	9	10.00	0.1	0.01	0.1
Sample_Depth	m	1	3.23	1	1	1
Cond	uS/cm	30	0.00		0	35
TDS	mg/l	30	0.00		568	1470
Sr	mg/l	23	0.00		464	1514
Mn_diss	mg/l	2	0.00		0.6	0.78
NH4	mg/l	14	12.50	0.025	0.025	0.6
Ba	mg/l	2	8.00	0.01	0.01	0.01
Zn_diss	ug/l	28	3.45	0	0	0.8
Cu_diss	ug/l	1	50.00	500	500	500
F	mg/l	1	3.70	500	500	500
SD4	mg/l	30	6.78	0.1	0.1	2.65
NO3	mg/l	30	3.23	1	1	792

PrintSaveRefreshClose

Select the [Save] button to export the data to a .TXT, .CSV or .XLS file.

5.3 Compare Sample

The **Compare Sample** report allows you to compare parameters for one sample to other samples in your database. This allows you to quickly determine the similarities and differences between your samples. When you select **Reports** from the main menu and then **Compare Sample**, the following dialogue will appear:




The top field in this dialogue “**Compare this sample to all active samples**” allows you to select a sample to be used in the report.




Press the **Select sample** button at the right side of this field and **Pick a Sample** dialogue will open up (as shown below). It allows you to select a sample for the Report.

ID	SAMPLEID	WATERTYPE	SAMPLE_DATE	ST
1	MW-1-92	Ca-CO3-SO4	8/15/1992	MV
2	MW-1-93	Ca-Na-SO4-Cl-HCO3	6/1/1993	MV
3	MW-1-94	Ca-Na-SO4-Cl	6/15/1994	MV
4	MW-1-95	Ca-Na-SO4-Cl-HCO3	7/30/1995	MV
5	MW-1-96	Ca-Na-SO4-Cl-HCO3	7/28/1996	MV
6	MW-1-97	Ca-Na-SO4-Cl-HCO3	6/15/1997	MV
7	MW-1-98	Na-SO4-Cl-HCO3	8/1/1998	MV
8	MW-3-92	Ca-SO4-HCO3	8/8/1992	MV
9	MW-3-93	Ca-HCO3-SO4	6/8/1993	MV
10	MW-3-94	Ca-SO4-HCO3	6/15/1994	MV
11	MW-3-95	Ca-HCO3-SO4	7/25/1995	MV

Double-click on the desired sample to load this into the Report options.

The **Compare Sample** report requires you to specify **Parameters** which will be used in the comparison analysis. To add new parameters, press the  button and select a parameter from the available list. You may also include parameter ratios, sums or differences (e.g. Na/Cl, Ca+Mg, Cl-SO4) in the **Parameters** field; simply type these

values in manually in the parameter field. To remove parameters from the list, press the  button. The default parameter settings for this report can be changed by modifying the parameters field and pressing the **[Set Default]** button in the lower left corner of this dialogue.

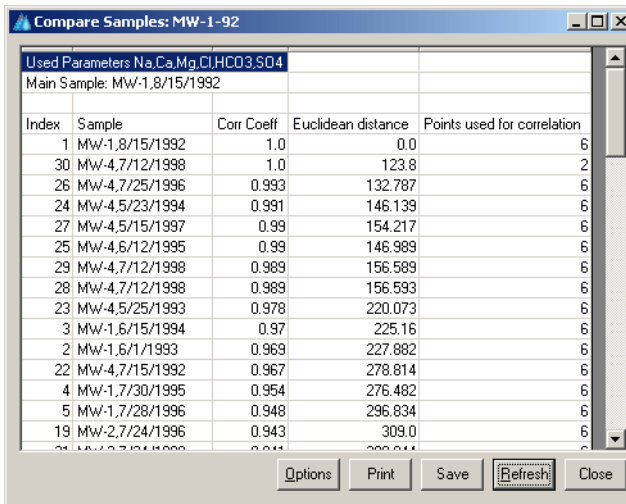
There are two options to select the samples from the active list:

- Select samples with correlation coefficient > #
- Use selected samples only

The **Select samples with correlation coefficient > #** option allows you to enter a correlation coefficient (r) value, and AquaChem will highlight (select) those samples in the active list once the comparison is completed.

When the **Use selected samples only** option is enabled, the Report will use only those samples that are selected (highlighted) in the active samples list, as part of the analysis.

Once you have specified the required options, press **[OK]** and the report will be generated. An example is seen below:

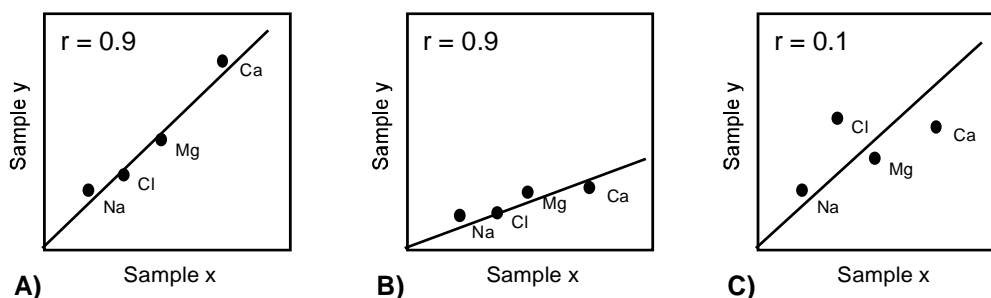


Index	Sample	Corr Coeff	Euclidean distance	Points used for correlation
1	MW-1,8/15/1992	1.0	0.0	6
30	MW-4,7/12/1998	1.0	123.8	2
26	MW-4,7/25/1996	0.993	132.787	6
24	MW-4,5/23/1994	0.991	146.139	6
27	MW-4,5/15/1997	0.99	154.217	6
25	MW-4,6/12/1995	0.99	146.989	6
29	MW-4,7/12/1998	0.989	156.589	6
28	MW-4,7/12/1998	0.989	156.593	6
23	MW-4,5/25/1993	0.978	220.073	6
3	MW-1,6/15/1994	0.97	225.16	6
2	MW-1,6/1/1993	0.969	227.882	6
22	MW-4,7/15/1992	0.967	278.814	6
4	MW-1,7/30/1995	0.954	276.482	6
5	MW-1,7/28/1996	0.948	296.834	6
19	MW-2,7/24/1996	0.943	309.0	6
21	MW-2,7/12/1998	0.941	309.0	6

The **Compare Sample** report uses a linear regression algorithm to generate the correlation coefficient and the Euclidean distance between a selected sample and all other active or selected samples. Samples having a chemical composition similar to the selected sample will have a correlation coefficient close to 1. The correlation coefficient is a function of ratios rather than absolute values. The difference in absolute concentrations is expressed by the Euclidean distance:

$$d_{ij} = \frac{\sum_{k=1}^n x_{ik} - x_{jk}}{n}$$

where x_{ik} denotes the k th variable measured on sample i and x_{jk} is the k th variable measured on sample j . For each sample, n variables are measured. The distance between sample i and sample j is d_{ij} .



In the examples above, comparing **Sample x** to **Sample y**, the following conclusions can be made:

- (A) high similarity
- (B) high similarity but sample y is diluted with respect to sample x
- (C) low similarity

5.4 Mix Samples

The **Mix Samples** Report generates solution concentrations resulting from the step-wise mixing of specified proportions of two selected samples from your database. When you select **Reports** from the main menu and then **Mix Samples**, the following dialogue will appear:

Mix Samples

Select Solutions

Mix: 18 OW-2-95 Na-Ca-Cl-SO4 7/21/1995 OW ...

With: 19 OW-2-96 Na-Ca-Cl-SO4 7/24/1996 OW ...

Parameters: Na, K, Ca, Mg

Mode: ☒ Simple Mixing ☐ Optimize

Simple Mixing

Add: 0.1 to 0.5 Parts of Solution 1 to Solution 2

Number of Steps: 5

Set Default OK Cancel

In the **Select Solutions** frame, press the button beside **Mix** and **With** fields to select a sample from the available solutions (samples) that are in your database. Each field contains a complete list of all active samples in your AquaChem database.

The **Parameters** list allows you to select the parameters that will be included in the mixing calculations. Typically you should select parameters that you know are common to each sample. (If you use parameters which are not common to both samples, then the report may not be successfully generated). To add a parameter to the list, press the button and select a parameter from the list. To remove a parameter, select the parameter then press the button. The default parameter settings for this report can be changed by modifying the parameters and pressing the **[Set Default]** button in the lower-left corner of the dialogue.

In the **Mode** frame there are two mixing options, the **Simple Mixing** mode and the **Optimize** mode. Each mode is described in detail below.

Simple Mixing Mode

When you select the **Simple Mixing** mode, AquaChem will mix the two selected samples in a step-wise process.

In the **Simple Mixing** frame, the **Add** field is the starting proportion (m) of Solution 1 in the mixture solution, while the **to** field is the final proportion (n) of Solution 1 in the mixture solution. The **Number of Steps** value is the number of uniform steps in which to get from (m) to (n) parts of Solution 1 in the mixture solution. Starting with (m) parts of Solution 1 and ($1.0 - m$) parts of Solution 2 in the mixture solution, AquaChem will

begin uniformly decreasing the proportion of Solution 1 in the mixture solution while maintaining a total proportion value of 1.0. The final mixture solution has (*n*) parts of Solution 1 and (1.0 - *n*) parts of Solution 2.

The figure below shows the **Sample Mixing Report**, and the results of mixing two solutions.

Sample Mixing Report								
Solution 1	MW-1.8/15/1992							
Solution 2	MW-2.8/1/1992							
Percentage of solution 1 in target solution 10%-50%								
Solution 1	1.0	0.10	0.20	0.30	0.40	0.50	0.0	
Solution 2	0.0	0.90	0.80	0.70	0.60	0.50	1.0	
Na	80.0	210.5	196.0	181.5	167.0	152.5	225.0	
K	1.5	1.14	1.18	1.22	1.26	1.3	1.1	
Ca	1000.0	1000.0	1000.0	1000.0	1000.0	1000.0	1000.0	
Mg	22.0	6.7	8.4	10.1	11.8	13.5	5.0	
Cl	125.0	269.9	253.8	237.7	221.6	205.5	286.0	
HCO3	125.0	138.77	137.24	135.71	134.18	132.65	140.3	
SO4	550.0	233.47	268.64	303.81	338.98	374.15	198.3	
	0.0	0.0	0.0	0.0	0.0	0.0	0.0	

The concentrations of each parameter in the first column are the concentrations for Solution 1. The concentrations of each parameter in the second column are the concentrations for a mixture of 0.1 parts Solution 1 with 0.9 parts Solution 2. The concentrations of each parameter in the third column are the concentrations for a mixture of 0.2 parts Solution 1 with 0.8 parts Solution 2, and so forth for the remaining columns.

Optimize Mode

In addition to simple mixing of two samples, AquaChem also allows you to determine the optimal mixing ratio of the two selected samples that most closely matches a target sample from the database. When you select the **Optimize** mode, you will see the following options (in the lower half of the Report options dialogue):

Optimize

Calculate the mixing proportions of solutions 1 and 2 that matches best the sample below

24 OW-4-94	Ca-SO4	5/23/1994	OW-4
------------	--------	-----------	------

Set Default OK Cancel

This mixing mode requires you to select a sample that you would like to match. AquaChem will mix the two initial samples in 2% increments until the Euclidean distance between the calculated mixture and specified resulting sample is minimized.

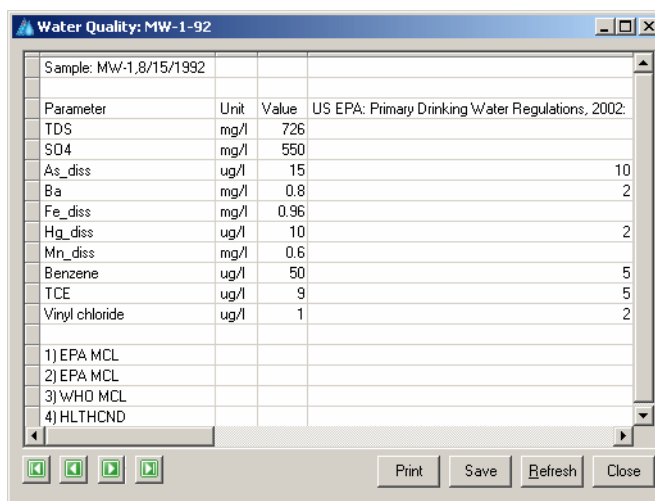
The **Optimize** mode can be used in a situation where you have three samples (ex. groundwater, seawater and freshwater) and you suspect that the groundwater is a mixture of the freshwater and the seawater. The **Optimize** mode can be used to estimate the percentage of freshwater and the percentage of seawater required to obtain the groundwater composition of selected parameters.

To generate a report with **Optimize** mode, select a sample from your database, and click [OK].

5.5 Water Quality Standards

The **Water Quality Standards** report provides a list of all parameters which exceed at least one of the established Guideline/Tolerance levels for the selected sample. If your sample contains no exceedences, then this Report will be blank.

When you select this report, a **Water Quality** window similar to the one below will appear:



Sample: MW-1,8/15/1992			
Parameter	Unit	Value	US EPA: Primary Drinking Water Regulations, 2002:
TDS	mg/l	726	
SO4	mg/l	550	
As_diss	ug/l	15	10
Ba	mg/l	0.8	2
Fe_diss	mg/l	0.96	
Hg_diss	ug/l	10	2
Mn_diss	mg/l	0.6	
Benzene	ug/l	50	5
TCE	ug/l	9	5
Vinyl chloride	ug/l	1	2
1) EPA MCL			
2) EPA MCL			
3) WHO MCL			
4) HLTHCND			

Each line of the report contains the **Parameter**, corresponding **Unit**, actual measured **Value** and guideline values (e.g. Maximum Contaminant level (**MCL**), Goal contaminant level (**GCL**), **Aesthetic Objectives (AO)**). Depending on the Guideline used, you may see one, two, or three guideline levels. The example above shows three levels, a maximum concentration limit, a tolerated concentration limit and a concentration goal. AquaChem allows the definition of multiple standards; the active

standard used for detecting exceedences in this report is defined in the **Preferences** dialogue available under the **File** menu.

The scroll arrows in the lower left corner of the **Report** window can be used to produce a report for other samples in your database.

5.6 Hardness Dependent Standards

The United States Environmental Protection Agency (EPA) has established standards for metals where toxicity is a function of hardness. The water quality criteria for metals can be expressed as “total recoverable” or “dissolved” for acute and chronic concentrations. The total recoverable also includes the sorbed concentration on particles. Acute standards offer protection from toxic effects from chemical concentrations during a shorter term exposure. Chronic standards offer protection from toxic effects from a chemical during long-term exposure.

EPA Freshwater standards are mainly used for hardness dependent standards, however similar standards can be found in different countries. The demo_basic database provided with AquaChem only includes the EPA Freshwater Standards, however you may create your own standards.

When you select this report, a **Freshwater Standards** window will appear (shown below):

Parameter	Unit	EPA MCL	WHO MCL	HLTHCND	Acute Total Recoverable	Acute Dissolved	Chronic Total Recoverable	Chronic Dissolved
Arsenic	ug/L	10	10		340	340	150	150
Cadmium	ug/L	5	3		2.133	2.014	0.2706	0.246
Chromium III	ug/L				1803.0	569.8	86.2	74.1
Copper	ug/L		1300	2000	14.0	13.44	9.33	8.96
Lead	ug/L			10	81.6	64.6	3.18	2.517
Mercury	ug/L				2.34	1.99	2.34	1.99
Nickel	ug/L		20		469.2	468.2	52.2	52.0
Silver	ug/L				3.784	3.217		
Zinc	ug/L			5000	119.8		2.66	2.62
Chromium VI	ug/L				16	15.7	11	10.58

Most Stringent

Options Export Close

The **Standards Tab** displays the non-hardness dependent standards, along with the acute and chronic standards for metals, according to the specified **Hardness (mg/L CaCO₃)**. When a new value is entered in the **Hardness** field, the acute and chronic values are automatically calculated and displayed in the table below for all metals that are hardness dependent.

The **Exceedences** tab shows a table with the measured hardness and measured parameter along with respective calculated standards based upon sample hardness for each sample in the active samples list.

Exceedences will be highlighted with different colors if acute or chronic standards are exceeded.

The **Function** tab shows the metal standards as a function of hardness value varying from 1-400 mg/L CaCO₃.

Creating a New Standard

By default, AquaChem provides the EPA Freshwater standard. However, if this standard is not suitable for your project, you may modify the standard or create a new standard from scratch.

To create a new hardness dependent standard, select the **Options** button, located in the bottom-right corner of the Fresh Water Standards dialog.

The **Hardness Dependent Standards** dialog will appear on your screen.

The screenshot shows the 'Hardness Dependent Standards' dialog box. It contains the following fields and controls:

- Standard:** EPA Freshwater
- Comment:** National Recommended Water Quality Criteria 2009, <http://www>
- Unit:** ug/L (dropdown menu)
- Hardness interval (mg/L CaCO₃):**
 - From:** 25
 - to:** 400
- Parameter:** Arsenic (dropdown menu)
- Type:** Constant (dropdown menu with add, delete, and refresh icons)
- Associated Total Recoverable Parameter:** As_tot (dropdown menu)
- Associated Dissolved Parameter:** As_diss (dropdown menu)
- CCC (chronic):** 150
- CMC (acute):** 340
- Conversion factor CCC:** 1
- Conversion factor CMC:** 1


At the bottom are buttons: Delete, Save, New, Duplicate, and Close.

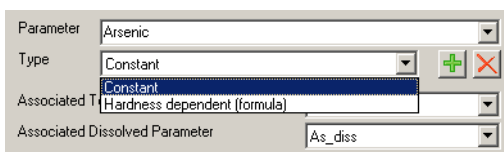
Information for the current standard will be shown in the Hardness Dependent Standard dialog. To create a new standard, click the **[New]** button.

Specify a **Name**, **Description** and **Units** for the new standard in the respective text fields.

Specify the desired **Hardness interval (mg/L CaCO₃)**. AquaChem will display the metal standards as a function of each hardness value within the specified range in the **Function** tab (of the Fresh Water Standards Dialog).

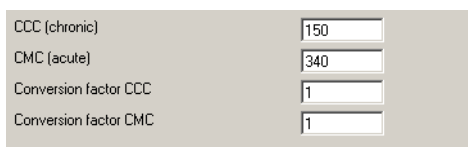
Adding Parameters to the Standard

- Click the  button, and then type the name of the parameter in the **Parameter** field.
- Select the desired type from the **Type** combo box. Choose **Constant** or **Hardness Dependent (formula)**.



Constant

The **Constant** option allows you to define a constant value for both the acute and chronic concentrations. In other words, these values will not change with changes in hardness.



The chronic concentration, also referred to as **Criteria Continuous Concentration (CCC)**, is an estimate of the highest concentration of a material in surface water to which an aquatic community can be exposed briefly without resulting in an unacceptable effect. The acute concentration, also referred to as **Criteria Maximum Concentration (CMC)**, is an estimate of the highest concentration of a material in surface water to which an aquatic community can be exposed briefly without resulting in an unacceptable effect.

You may also specify the conversion factor for both CCC and CMC. The term “Conversion Factor” is the recommended conversion factor for converting a metal criterion expressed as the total recoverable fraction in the water column to a criterion expressed as dissolved fraction in the water column.

Hardness Dependent (Formula)

This options allows you to express the freshwater criterion for the metal as a function of hardness (mg/L) in the water column. The hardness level may be specified in the main **Fresh Water Standards** dialog, in the **Standards** tab (shown below).

When the **Hardness Dependent (Formula)** option is selected, the following options will appear:

The criteria values for CMC and CCC are calculated using the following formulas:

$$\text{CMC} = \exp\{m_A[\ln(\text{hardness})] + b_A\}$$

$$\text{CCC} = \exp\{m_C[\ln(\text{hardness})] + b_C\}$$

Where m_A , b_A , m_C and b_C are empirical coefficients for the metal.

To express the metal criterion as dissolved fraction in the water column, a conversion factor is applied to the above formulas. The following formulas are used to define the conversion factor as a function of hardness.

$$\text{CF}(\text{CMC}) = a - [\ln(\text{hardness}) * b]$$

$$\text{CF}(\text{CCC}) = c - [\ln(\text{hardness}) * d]$$

To view a table of parameters used for calculating freshwater dissolved metals criteria that are hardness-dependent, please refer to the “National Recommended Water Quality Criteria” document by the United States Environmental Protection Agency, available on-line at the following website:

<http://epa.gov/ost/criteria/wqctable/>

Saving Standards

To save the changes made to your hardness dependent standard, click the **[Save]** button located at the bottom of the dialog.

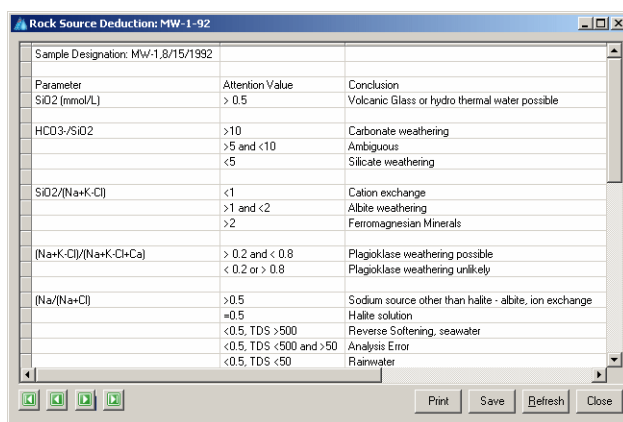
Duplicating Standards

You may wish to create a duplicate of an existing standard to use as a template, for creating new standards. To create a duplicate, click the **[Duplicate]** button located at the bottom of the dialog.

5.7 Rock Source Deduction

The **Rock Source Deduction** report allows you to gain insight into the possible origin of the water sample. The results are a general overview based on ion ratios found in a sample which are compared to ratios of the respective ions in reactive minerals. If results do not meet expectations, the result should be confirmed with more detailed study based on multiple samples, aquifer mineralogy analysis, modeling and with the use of plots.

When you select this report, the following **Rock Source Deduction** window will appear:



Parameter	Attention Value	Conclusion
SiO2 (mmol/L)	> 0.5	Volcanic Glass or hydro thermal water possible
HCO3-/SiO2	>10	Carbonate weathering
	>5 and <10	Ambiguous
	<5	Silicate weathering
SiO2/(Na+K+Cl)	<1	Cation exchange
	>1 and <2	Albite weathering
	>2	Ferromagnesian Minerals
(Na+K+Cl)/(Na+K+Cl+Ca)	> 0.2 and < 0.8	Plagioclase weathering possible
	< 0.2 or > 0.8	Plagioclase weathering unlikely
[Na]/(Na+Cl)	>0.5	Sodium source other than halite - albite, ion exchange
	=0.5	Halite solution
	<0.5, TDS >500	Reverse Softening, seawater
	<0.5, TDS <500 and >50	Analysis Error
	<0.5, TDS <50	Rainwater

The following table provides a summary of the criterion for the Rock Source Deduction Report.

Parameter	Attention Value	Conclusion	Comments
SiO2 (mmol)	>0.5	Volcanic glass or hydrothermal water possible	The solubility of quartz at 20° is X mg/L.

HCO ₃ ⁻ /SiO ₂	>10 >5 and <10 <5	Carbonate weathering Ambiguous Silicate weathering	Low carbonate SiO ₂ ratios indicate, that these ions are released from silicate minerals, especially if TDS is low.
SiO ₂ /(Na+K-Cl)	<1 >1 and <2 >2	Cation exchange Albite weathering Ferromagnesian minerals	
(Na+K-Cl)/ (Na+K-Cl+Ca)	0.2 - 0.8 <0.2 or >0.8	Plagioklas weathering possible Plagioklas weathering unlikely	
Na/(Na+Cl)	>0.5 =0.5 <0.5 TDS >500 <0.5 TDS <500 and >50 <0.5 TDS <50	Sodium Source other than halite - albite, ion exchange Halite Solution Reverse Softening, seawater Analysis Error Rainwater	
Mg/(Ca+Mg)	=0.5 and (HCO ₃ /Si)>10 <0.5 >0.5 <0.5 and (HCO ₃ /Si)<5 >0.5	Dolomite Weathering Limestone-Dolomite Weathering Dolomite Dissolution, calcite precipitation, or seawater Ferromagnesian Minerals Granitic weathering	
Ca/(Ca+SO ₄)	=0.5 <0.5 and pH<5.5 <0.5 and pH neutral >0.5	Gypsum dissolution Pyrite oxidation Calcium removal - ion exchange or calcite precipitation Calcium source other than gypsum - carbonate or silicates	
TDS	>500 <500	Carbonate weathering or brine or seawater Silicate weathering	
Cl/Sum Anions	>0.8 and TDS>500 >0.8 and TDS<100 <0.8	Seawater or brine or evaporites Rainwater Rock weathering	
HCO ₃ / Sum Anions	>0.8 >0.8 and SO ₄ >20meq/L <0.8 and sulfate low	Silicate or carbonate weathering Gypsum dissolution Seawater or brine	
Calcite Saturation Index	>0 =0 <0	Oversaturated with respect to Calcite Saturated with respect to Calcite Undersaturated with respect to Calcite	

The report will complete the **Parameter** analysis, check the **Attention Values**, and produce a result value and **Conclusion** on the rock source.

If the report values shows “n/a”, it means that no data is available and the calculation could not be performed.

The scroll arrows in the lower right corner of the **Rock Source Deduction** Report window can be used to produce a report for other samples in your database.

For more details please see the reference below:

Hounslow, A.W. 1995. *Water Quality Data – Analysis and Interpretation*. CRC Press LLC. p.85.

5.8 Statistics

AquaChem comprises most of the commonly used statistical methods and techniques for the analysis of water samples:

- Summary Statistics
- Correlation Matrix
- Trend Analysis
- Outlier tests, and
- Tests for Normality
- Alert Levels

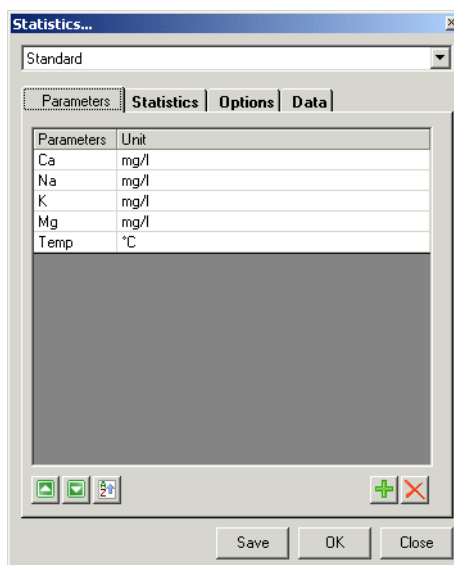
These analyses are available under **Reports>Statistics** and are discussed in detail below.

The statistics used in AquaChem were developed in accordance with USEPA guidelines outlined in the following document:





Guidance for Data Quality Assessment, Practical Methods for Data Analysis, EPAQA/G-9 QA00 Update. July 2000. USEPA Office of Environmental Information, Washington

5.8.1 Summary Statistics

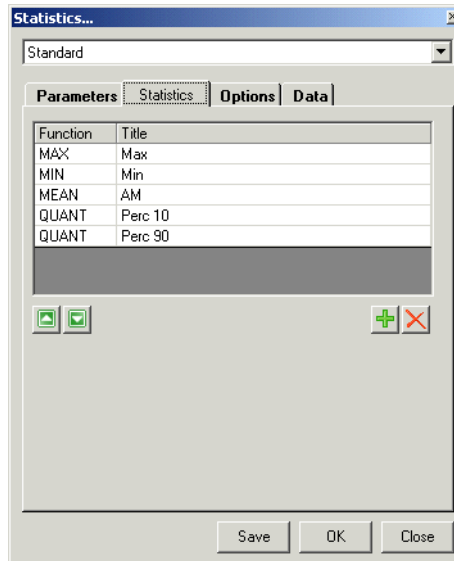
The **Summary Statistics** report calculates basic statistics for all active samples in your database. When you select **Reports** from the main menu, then **Statistics**, and, finally, **Summary Statistics** from the sub-menu, the following **Statistics** options dialogue will appear:




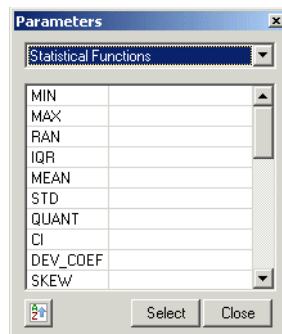
The above dialogue allows you to customize your summary statistics report using the available tabs. At the top of this dialogue is a combo box which allows you to choose a previously saved combination of settings.




In the first tab, **Parameters**, you must specify the parameters for which the summary statistics are to be calculated. Click the  button to produce a standard parameter selection dialogue and choose the desired parameters by highlighting them and clicking **[Select]** button. When finished with parameter selection click **[Close]**. You can change the order of the chosen parameters by clicking the  and  buttons. Delete any parameter by clicking the  button.

Next, click on the **Statistics** tab.

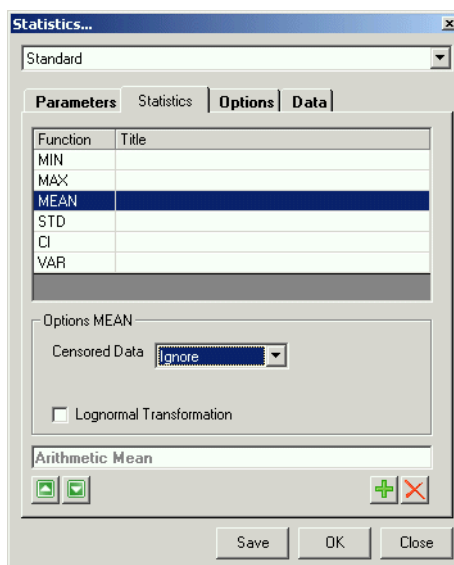


Use this dialogue to select the statistics for your data. Click the  button to produce the following dialogue:



Scroll through the available list of statistics to select the desired ones. Click **[Select]** to add a chosen statistic. When finished, click **[Close]**. The statistic abbreviation will be listed in the **Function** column of the statistic list. The **Title** column lists the full statistic names and these names will appear as headers on the output. You can modify these entries by double-clicking in the cell and entering a new **Title**. You can change the order of the chosen statistics by clicking the  and  buttons. Delete any parameter by clicking the  button.

Once the statistics are chosen, you can modify some of their options in the **Summary Statistics** dialogue. To access statistic options simply highlight it in the list, and available options will be displayed in the bottom portion of the dialogue:



AquaChem allows you to decide how to deal with censored data. The **Censored Data** combo box allows you to select the method used to manage non-detects. The options are Auto, Ignore, Multiply, Constant, Trimmed, Winsorize. Depending on the method chosen you may have to enter additional data (e.g. if you chose Multiply, you will have to enter the Factor to multiply by). Choosing Auto will subject the censored data to the method specified in the Preferences (for more details see “QC Metadata” on page 122 in Chapter 3).

The following is a description of each Statistical Analysis available for this report:

MIN (minimum): lowest recorded value for the parameter

MAX (maximum): highest recorded value for the parameter

RAN (range): maximum minus minimum

IQR (interquartile range): Q75 minus Q25

MEAN (Arithmetic mean): average value for the parameter

STD (Standard Deviation): square root of the variance

$$s = \sqrt{s^2}$$

QUANT (non parametric quantile): The Quantile is a data value, that is greater than or equal to a given fraction of the data values. If x is the p quantile, then at least the fraction p of the value set lie at or below x , at least $1-p$ of the value lie at or above x . For the non parametric quantile calculation no assumption is made for the underlying data set. The data is ranked from the lowest to the highest value. The quantile p is then calculated depending as follows

if number of points is odd:

$$\tilde{X} = X_{([n+1]/2)}$$

if number of points is even:

$$\tilde{X} = \frac{X_{(n/2)} + X_{([n/2]+1)}}{2}$$

CI (confidence interval): The confidence interval gives an estimated range of values which is likely to include the sample mean. The estimated range is calculated from a given set of sample data. Confidence intervals are usually calculated so that this percentage is 95%, but AquaChem can produce 90%, 99%, 99.9% (or any other) confidence intervals for the mean. The confidence range is calculated as the upper minus the lower confidence interval

$$\left(\bar{x} + t_{1-\alpha, n-1} \cdot \frac{s}{\sqrt{n}} \right) - \left(\bar{x} - t_{1-\alpha, n-1} \cdot \frac{s}{\sqrt{n}} \right)$$

DEV_COEF (Deviation Coefficient): Coefficient of Variance calculated using the formula below.

$$CV = \frac{s}{\bar{x}} \cdot 100\%$$

SKEW (skewness): skewness is calculated according to the following formula:

$$skew = \frac{n}{(n-1)(n-2)} \sum \left(\frac{x_i - \bar{x}}{s} \right)^3$$

KURT (kurtosis): kurtosis is calculated according to the following formula:

$$kurt = \frac{\sum_{i=1}^n (x_i - \bar{x})^4}{n \cdot s^4} - 3$$

N (number of samples): number of samples that have a measured value for the selected parameter.

N_ND (number of non-detects): number of samples that list the value for the selected parameter as a non-detect

PERC_ND (percent of non-detects): number of non-detects divided by the total number of samples and multiplied by one hundred

VAR (Variance): average square distance from points to sample mean

$$s^2 = \frac{1}{n-1} \sum_{i=1}^n (y_i - \bar{y})^2$$

QUANT_PAR (parametric quantile): For the parametric quantile calculation the data is assumed to be normally distributed. A 0.25 quantile thus corresponds to the Z value of the normal distribution for a probability of 0.25.

N_EXC (number of exceedences): number of samples that have exceedences for the selected parameter

PERC_EXC (percent of exceedences): number of exceedences for the parameter divided by the total number of samples and multiplied by one hundred

N_D (number of detects): number of samples for which the selected parameter could be detected

PERC_D (percent of detects): number of detects for the parameter divided by the total number of samples and multiplied by one hundred

N_DBS (number of detects below standard): number of values below standard for the selected parameter

PERC_DBS (percent of detects below standard): number of detects below standard divided by the total number of samples and multiplied by one hundred

N_DAS (number of detects above standard): number of values above standard for the selected parameter

PERC_DAS (percent of detects above standard): number of values above standard for the selected parameter divided by the total number and multiplied by one hundred

TSTAT (t-statistic): calculates t-statistic for the parameter

LCL (lower confidence limit): calculates lower confidence limit

$$\bar{x} - t_{1-\alpha, n-1} \cdot \frac{s}{\sqrt{n}}$$

UCL (upper confidence limit): calculates upper confidence limit

$$\bar{x} + t_{1-\alpha, n-1} \cdot \frac{s}{\sqrt{n}}$$

TI (tolerance interval): The tolerance interval is a way of determining a range that (with a certain confidence level) will contain a certain percentage of the population

LTl (lower tolerance limit): calculates lower tolerance limit

UTl (upper tolerance limit): calculates upper tolerance limit

MKS (Mann Kendall Statistic): indicates increasing or decreasing trend

MKZ: (Mann Kendall Statistic): approximates Z value for calculating probability

GEOMEAN (Geometric Mean): the mean of n numbers expressed as the n-th root of their product.

P_EXC: Probability of exceeding a given value. This function assumes that the data is normally distributed. It calculates the standard deviation and average for a given parameter, and then calculates the probability that the specified threshold is exceeded. Typically the threshold is defined as the water standard for the given parameter. The P_EXC function then returns the probability that exceeding values will be encountered.

LPR/UPL: The prediction interval estimates future values based on present or past background samples. Prediction intervals tend to be applied in detection monitoring in two main ways. They can be used to compare compliance wells with background wells, or they can be used for intrawell comparisons of monitoring wells. When comparing compliance wells to a background well, if the compliance wells come from the same uncontaminated water source, the upper prediction limit should be greater than or equal to the data collected from compliance wells. Compliance data greater than upper prediction limits is

indicative of contamination. For intrawell comparisons, a range of values is determined which future values collected from the same well should fall within. Any data collected in the future which does not fall within that specified range is an indication that a once uncontaminated water supply is now contaminated. Below is the formula for calculating lower and upper prediction limits:

$$\bar{x} \pm ts \sqrt{1 + \frac{1}{n}}$$

where,

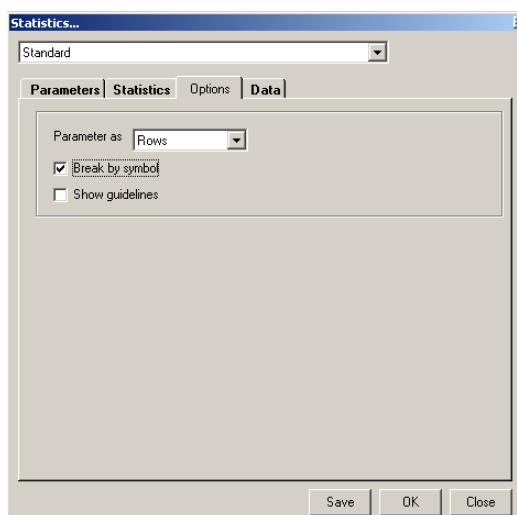
\bar{x} = mean

$t(n-1, \alpha)$ = t-statistics value (function of confidence and number of points)

s = standard deviation

n = number of data points

The next tab allows you to specify the output settings:

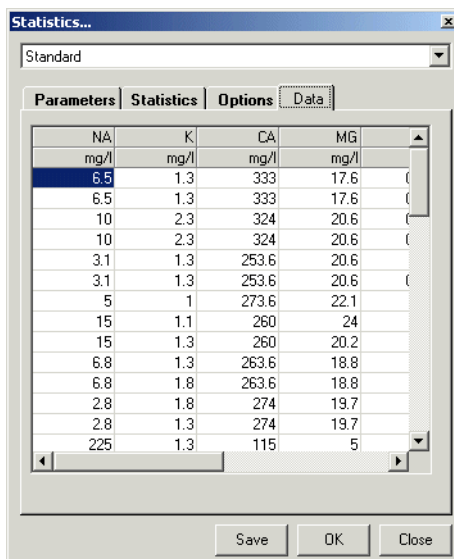


The **Parameter** as combo box allows you to set the parameter orientation. When **Rows** is selected, the parameters will be displayed in rows and statistics in columns. When **Columns** is selected, the orientations are reversed with parameters displayed in columns and statistics in rows.

AquaChem allows you to break the **Summary Statistics** output by symbol. When this option is selected, the statistics will be calculated and displayed for every symbol.

Thus, if you have a symbol assigned to every sample, all samples with the same symbols will be grouped and statistics performed on those groups.

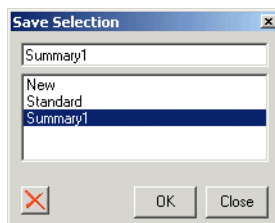
The **Data** tab provides you with a preview of data of the selected parameters:




The **Statistics...** dialog box is shown with the **Data** tab selected. The table displays data for four parameters: NA, K, CA, and MG, all measured in mg/l. The data is organized into rows, with the first row highlighted in blue.

NA	K	CA	MG
mg/l	mg/l	mg/l	mg/l
6.5	1.3	333	17.6
6.5	1.3	333	17.6
10	2.3	324	20.6
10	2.3	324	20.6
3.1	1.3	253.6	20.6
3.1	1.3	253.6	20.6
5	1	273.6	22.1
15	1.1	260	24
15	1.3	260	20.2
6.8	1.3	263.6	18.8
6.8	1.8	263.6	18.8
2.8	1.8	274	19.7
2.8	1.3	274	19.7
225	1.3	115	5

If you are satisfied with the settings for the **Summary Statistics** report, before generating it, you may wish to save these settings by clicking **[Save]** button. The following **Save** dialogue will be loaded:



The **Save Selection** dialog box is shown. It contains a text field with 'Summary1' and a list box with 'New', 'Standard', and 'Summary1'. The 'Summary1' option in the list box is selected. There are buttons for 'OK', 'Close', and a red 'X' button to delete the settings.

Type in a new name, or select an already existing name if you wish to update the settings. This dialogue also allows you to delete the settings that are no longer needed. Simply select the name of the settings you wish to delete and click the  button.

Click **[OK]** to save the settings.

Click **[OK]** to generate the report.



An example of a finished report is displayed below:

Parameter	Unit	EPA MCL	EPA MCL	WHO MCL	HLTHCND	Minimum	Maximum	Arithmetic Mean	Standard deviation
pH_field						0.6	62	7.63	6.6
Cond	uS/cm					250	1470	734.5	410.3
TDS	mg/l				500	140	1514	642.9	465.3
Ca	mg/l					3	22010	801.1	2563.2
Mg	mg/l					0.04	3800	115.4	509.0
Na	mg/l				200	2.8	38400	914.5	4542.7
K	mg/l					1	13400	238.0	1573.4
Cl	mg/l				250	2.8	218000	6078.4	30171.2
HCO3	mg/l					1.2	4130	283.7	491.8
SO4	mg/l				500	1	62400	1408.8	8887.6
F	mg/l	4	4	1.5		0.1	2.65	0.78	0.866
NO3	mg/l	10	10	50		0.5	4	0.913	1.344

You may print this report, or save it in .XLS, .TXT, or .CSV format.

5.8.2 Correlation Matrix

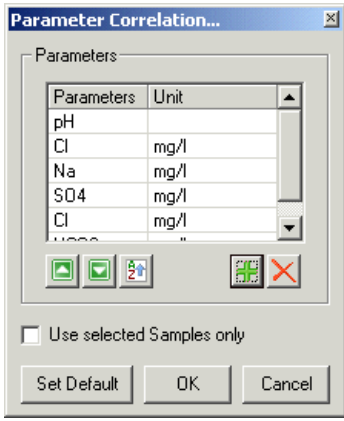
The **Correlation Matrix** Report shows a simple correlation matrix of the parameters for the samples in your database. This allows you to quickly determine the similarities or differences between your samples. When you select **Reports** from the main menu and then **Statistics>Correlation Matrix**, the following dialogue will appear:

The **Correlation Matrix** Report requires you to specify parameters which will be used in the correlation analysis. To add new parameters, press the  button and select a parameter from the available list. You can also include parameter ratios, sums or differences (e.g. Na/Cl, Ca+Mg, Cl-SO4) in the **Parameters** field; simply type these values in manually in the parameter field. To remove parameters from the list, press the  button.

When the **Use Selected samples only** option is enabled, only the samples which are selected in the active list will be used. If this option is disabled, then all samples in the active list will be used.

The **Unit** field beside every parameter contains a combo box listing the available concentration units for the correlation of the chemical parameters. Units of mol/l are often more suitable for correlating than mg/L, because it gives you an idea as to which minerals have been dissolved.

Once you have specified the required options, press **[OK]** and the report will be generated as shown in the figure below:



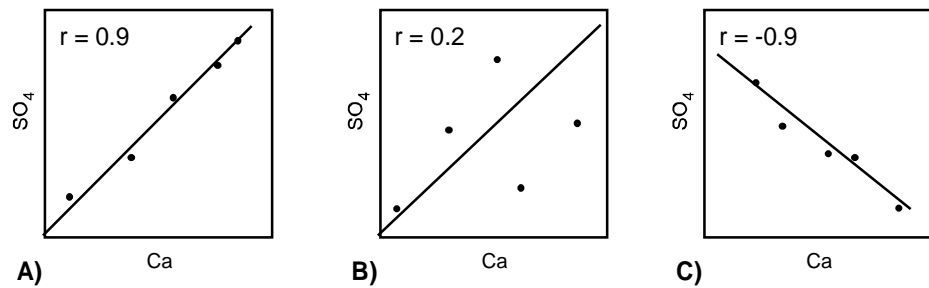
The dialog box titled "Parameter Correlation..." contains a list of parameters with their units. The list includes pH, Cl (mg/l), Na (mg/l), SO4 (mg/l), and Cl (mg/l). Below the list are buttons for adding, removing, and refreshing the list. There is a checkbox labeled "Use selected Samples only" which is currently unchecked. At the bottom are buttons for "Set Default", "OK", and "Cancel".

Correlation...					
Total Number of Samples: 79					
Correlation coefficient					
	pH_field	Ca	Cl	SO4	
pH_field	1.0	-8.3E-2	-0.511	-0.17	
Ca mg/l		1.0	0.475	-1.5E-2	
Cl mg/l			1.0	0.895	
SO4 mg/l				1.0	
y axis intersection					
	pH_field	Ca	Cl	SO4	
pH_field	0.0	1.1E+3	6.4E+4	3.7E+3	
Ca mg/l		0.0	1.7E+3	2.1E+3	
Cl mg/l			0.0	229.76	
SO4 mg/l				0.0	
Slope of regression line					
	pH_field	Ca	Cl	SO4	
pH_field	1.0	-32.09	-8.5E+3	-187.85	
Ca mg/l		1.0	5.38	-4.3E-2	
Cl mg/l			1.0	0.219	
SO4 mg/l				1.0	
Number of datapoints for regression					
	pH_field	Ca	Cl	SO4	
pH_field	75	70	68	47	
Ca mg/l		74	68	51	
Cl mg/l			68	46	
SO4 mg/l				51	
Spearman's Rank Correlation Coefficient					
	pH_field	Ca	Cl	SO4	
pH_field	1.0	-0.525	-0.396	-0.416	
Ca mg/l		1.0	0.383	0.471	
Cl mg/l			1.0	1.5E-2	
SO4 mg/l				1.0	
Options Print Save Refresh Close					

The **Correlation Matrix** report generates a correlation matrix for a specified number of sample parameters that are common to all samples. A linear regression routine calculates the regression coefficient (r), and the slope and intercept of the regression line.

$$r = \frac{\sum_{i=1}^n x_i y_i - \frac{\sum_{i=1}^n x_i \sum_{i=1}^n y_i}{n}}{\sqrt{\left[\sum_{i=1}^n x_i^2 - \frac{(\sum_{i=1}^n x_i)^2}{n} \right]} \sqrt{\left[\sum_{i=1}^n y_i^2 - \frac{(\sum_{i=1}^n y_i)^2}{n} \right]}}$$

The figure below shows the correlation of **Ca** against **SO4** for three different scenarios.



The correlation of Ca and SO₄ in three samples above is as follows:

- (A) high positive correlation
- (B) low positive correlation
- (C) high negative correlation

It is often useful to check the correlation results graphically in a scatter plot to easily identify outliers which might influence the correlation result. The **Scatter** plot is available as one of the plot types, under the **Plots** menu. In the **Scatter** plot options, access the **Edit Lines** option under the **Symbols** tab. In this dialogue, you can calculate the regression curve. For more details, see Scatter Plot in Chapter 4.

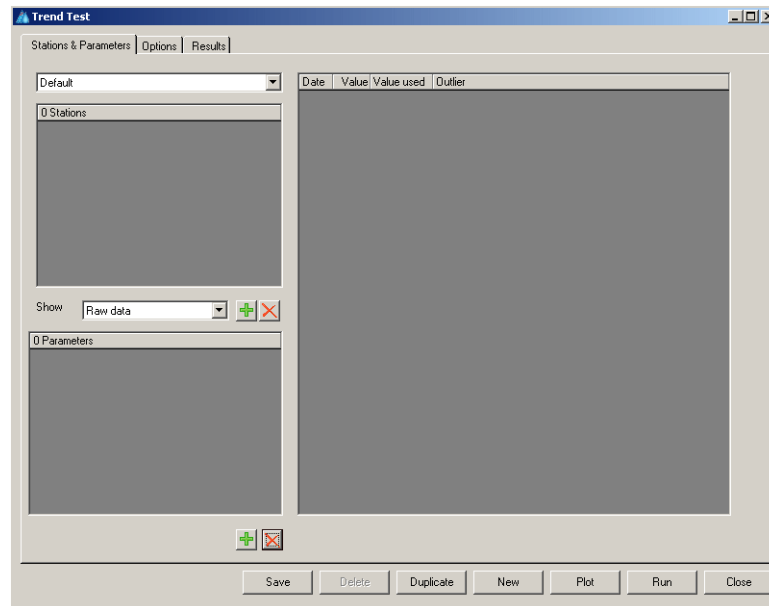
The Correlation Matrix also includes the Spearman Rank Correlation coefficient. This value is calculated using the assigned rank, rather than exact values. This coefficient is useful when it is not economically feasible to obtain the exact values. Spearman Rank Correlation coefficient is a good indicator of a relationship between two variables, in case this relationship is not linear.

5.8.3 Trend Analysis

The **Trend Analysis** option estimates a potential positive or negative trend that may be present in the data. A trend analysis may be interesting for three types of problems:

- When detecting an increasing trend becomes important to show that the contaminants in the down gradient wells increase with time and may eventually exceed its water quality standards.
- For compliance monitoring projects, the success of groundwater cleanup and remediation works can be verified by testing if the contaminant shows a decreasing trend.
- Finally, if a dataset is to be used in a test assumption of independent data, trend analysis can be used in order to ensure that this data does not include any trend at all.

When you select **Reports>Statistics>Trend Analysis**, the following dialogue will appear:



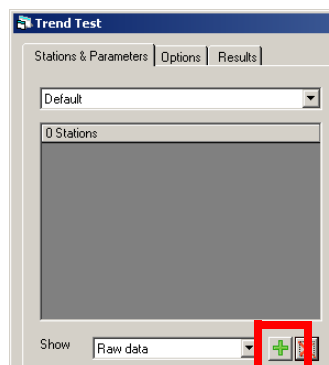
The **Trend Test** dialog consists of three tabs: **Stations & Parameters**, **Options** and **Results**. Each tab is described below.

Stations & Parameters

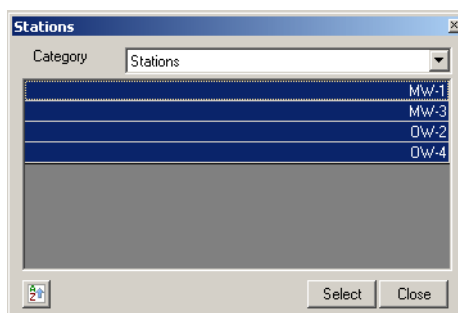
This tab allows you to specify which stations and parameters to include in the trend analysis.

Adding Stations

To add one or more stations, select the  button, located beside the **Show** combo box.

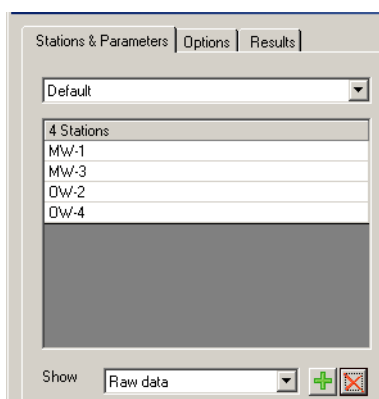


The **Stations** dialog will appear on your screen. The stations dialog lists all stations currently displayed in the **Active Stations** list. **Note:** If your database consists of many stations, it may be desirable to perform a query (see “Find” on page 136) such that only those stations required for the trend analysis are displayed in the **Active Stations** list.




Select the desired stations from the **Stations** dialog, and then select the **[Select]** button. Click the **[Close]** button to close the dialog.

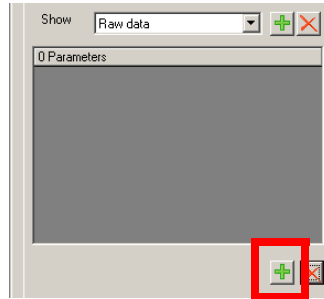
The selected stations will then be added to the **Stations** list, in the **Trend Test** dialog.



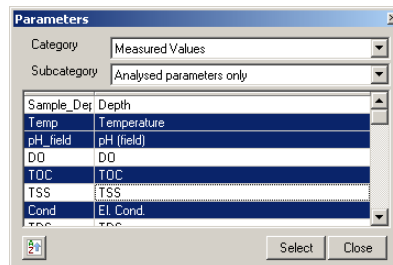
Selecting Parameters

The AquaChem trend analysis report allows you to analyze any combination of stations and parameters in a single run.

To add one or more parameters to the analysis, select the  button, located below the **Parameters** list.



The **Parameters** dialog will appear on your screen where you can select one or more parameters for the analysis. Please note that the list of parameters in the dataset may be very extensive, and you may not want to include parameters which have not been analyzed in any of the samples in the current selection of stations. Select the subcategory filter “Analyzed parameters only” to show only those parameters that have been analyze.

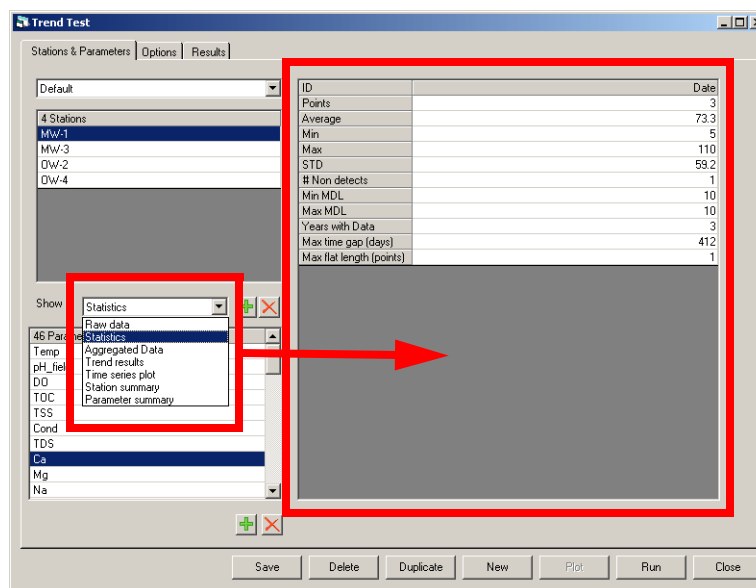


Select the desired parameters, and then select the **[Select]** button. Click the **Close** button to close the dialog.

Exploring the Dataset

Before you run the actual trend analysis, with potentially hundreds of stations and parameters, you should first explore the dataset and check whether there are stations or parameters which should be excluded from the analysis due to insufficient data.

Use the **Show** combo box, to display various information in the adjacent display area about the selected station and parameter



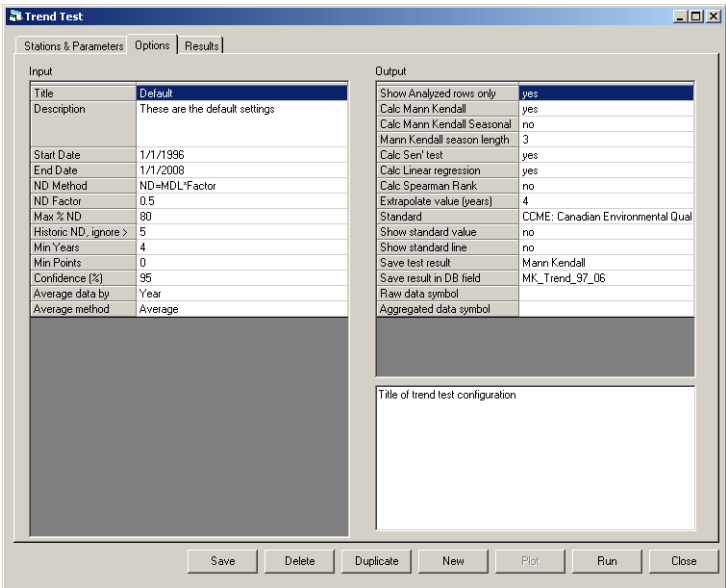
The show combo box provides the following options:

Raw Data	Shows all data points including samples marked as outliers. Note that outliers will not appear in other previews, e.g., in plots
Statistics	Provides a short summary of statistics for specified dataset.
Aggregated Data	Shows the average measurement in each aggregation interval. For example, if you aggregate over 12 months, you will see a yearly averaged sample. Please note that the concentration as well as the time is averaged. If all measurements were taken in Q4, the averaged point will also plot in Q4, not in the middle of the year.
Trend Test	Summarizes the trend test results for the current parameter and station selection. The information of each of the 5 tests are shown: Man Kendall, seasonal Mann Kendall, Sen Test, Linear Regression and Spearman Correlation. Please note that the linear regression coefficient detects linear relations only and is sensitive to outliers. The Spearman's rank correlation coefficient detects any monotonic trend and is not limited to linear.
Time Series Plots	Generates a time series diagram including the Sen slope, the linear regression and the water standard for the active parameter. The data points are shown as raw data and averaged data points if this option has been selected.
Station Summary	Displays a summary of each station in the database that has an analyzed parameter value, including the Min, Max, Count, date of the First Sample and date of the Last sample.

Parameter Summary	Displays a list of stations that contain data for the selected parameter, including the Min, Max, Count, date of the First Sample and date of the Last sample.
--------------------------	--

Options

The options tab allows you to configure the various input and output options for the trend analysis. Once you have specified the desired options, you can save the configuration by selecting the **[Save]** button from the row of buttons along the bottom of the dialog.



Input

The input options allow you to configure various input parameters for the trend analysis. Each option is described briefly below:

Title	The title of the trend test configuration
Description	A description of the trend test configuration
Start Date	Start data for trend test analysis. Samples before this date will be ignored in the analysis. Format: MM/DD/YYYY
End Date	End data for trend test analysis. Samples after this date will be ignored in the analysis. Format: MM/DD/YYYY
ND Method	Method for handling non detect values

ND Factor	The factor by which all non detect concentration are multiplied or the constant value that is assigned to all non detect concentrations.
MAX % ND	The maximum percentage of non detects allowed in dataset. If this threshold is exceeded, no test will be performed.
Historic ND, ignore >	For long time intervals the method detection limit may have changed between early and recent data, and more recent detection limits are often lower. When multiplying non detects by a factor, the dataset may become biased. Chose a ratio of historic/modern MDL above which you decide to ignore non detects. For example, if the value is set to 10 and the modern (minimum) MDL is 0.1, all non detects greater than < 1 will be ignored.
Min Years	The minimum number of years with at least 1 sample in the test interval. If this threshold is not reached, no test will be performed.
Min Points	The minimum number of points allowed in the dataset. If this threshold is not reached, a test will not be performed.
Confidence (%)	Confidence for trend tests.
Average data by	Choose the desired time interval for averaging data or disable averaging.
Average method	Select the desired averaging method that will be used to average data within the specified time interval.

Output

The output options allow you to configure how the results of the trend analysis are displayed in the Result tab and time series plots. The output options are described briefly below.

Show Analyzed Rows only	Only display rows where the data quality criteria have been matched. This is useful for tests with a large number of stations and parameters where many records do not reach the data quality criteria, e.g., there are not enough points to too many non detects.
Calc. Mann Kendall	Show/hide the results of the Mann Kendall test
Calc Mann Kendall Seasonal	Show/hide the results of the Seasonal Mann Kendall test
Mann Kendall season length	Specify the length of season for the Mann Kendall test, specified in months.
Calc. Sen' Test	Show/hide the results of the Sen test
Calc Linear Regression	Show/hide the results of the linear regression

Calc. Spearman Rank	Show/hide the results of the Spearman Rank correlation
Extrapolate Value (years)	Define a prediction year in number of years after the end date for test data.
Standard	Specify the standard that will display in the results table
Show standard value	Show/hide the water quality standard in the table results.
Show standard line	Select whether you want to display the water quality standard as a line on the time series diagrams.
Save test result	<p>Specify the test result that is saved to the aquachem database or select none if the results should not be saved to the database. This option requires that a meta data parameter has previously been defined to hold the test result.</p> <p>If this option is selected, a value of 1 will be stored for every parameter/station combination where a statistical trend was detected. This can be useful if you would later query for test results to detect patterns that may explain the occurrence of outliers.</p>
Save result in DB field	Select which parameter meta data field should be used to store the information. If you do not find an appropriate field, you may create it in the File > Database utility.
Raw data symbol	Define the desired symbology for the raw data points on the time series plots.
Aggregated data symbol	Define the desired symbology for the aggregated data points on the time series plot.

Results

Once you have specified the Stations & Parameters and have configured the input and out Options, you can run the trend analysis. To run the trend analysis, click the **[Run]** button located along the bottom of the **Trend Test** dialog.

Depending on how many parameters and stations are include in the test, this may take a few minutes to complete. The results will display in the **Results** tab.

Trend Test

Stations & Parameters

Options

Results

Settings

Default

Z(a=0.05)

1.644954

Start date

1/1/1952

End date

1/1/2008

Min Years

4

Data aggregation

Average

Min Points

8

Max ND % allowed

80

Correction for ND method

ND=MDL*Factor

ND Factor

0.5

Row	Station	Parameter	Unit	Standard	min MDL	Non Detects	%	Outliers	%	MDL>0%	%	Total samples	Valid samples	Years with Data	D
1	MW-1	Sample_Depth		0	0	0	0	0	0	0	0	7	7	100	7 p
2	MW-1	Temp	°C	0	0	0	0	0	0	0	0	7	7	100	7 p
3	MW-1	pH_Field		0	0	0	0	0	0	0	0	7	7	100	7 p
4	MW-1	Cond	uS/cm	0	0	0	0	0	0	0	0	7	7	100	7 p
5	MW-1	TDSS	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
6	MW-1	Ca	mg/l	0	10	1	14	0	0	0	0	7	7	100	7 p
7	MW-1	Mg	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
8	MW-1	Na	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
9	MW-1	K	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
10	MW-1	Cl	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
11	MW-1	HCO3	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
12	MW-1	SO4	mg/l	0	0	0	0	0	0	0	0	7	7	100	7 p
13	MW-1	As_diss	ug/l	0	0.01	1	14	0	0	0	0	7	7	100	7 p
14	MW-1	Ba	mg/l	2	0	0	0	0	0	0	0	7	7	100	7 p
15	MW-1	Fe_diss	mg/l	0	0	0	0	0	0	0	0	5	5	100	5 p
16	MW-1	Hg_diss	ug/l	2	0.001	4	57	0	0	0	0	7	7	100	7 p
17	MW-1	Mn_diss	mg/l	0	0	0	0	0	0	0	0	4	4	100	4 p
18	MW-1	Pb_diss	ug/l	0	0.007	4	67	0	0	1	17	7	6	86	6 p
19	MW-1	Boreon	ug/l	0	1	2	28	0	0	0	0	7	7	100	7 p
20	MW-1	TCE	ug/l	0	0	0	0	0	0	0	0	7	7	100	7 p
21	MW-1	PCE	ug/l	0	2	4	57	0	0	0	0	7	7	100	7 p
22	MW-1	14C	% mod	0	0	0	0	0	0	0	0	5	5	100	5 p
23	MW-1	190		0	0	0	0	0	0	0	0	7	7	100	7 p
24	MW-1	2H		0	0	0	0	0	0	0	0	7	7	100	7 p
25	MW-1	Tribum	T.U.	0	0	0	0	0	0	0	0	7	7	100	7 p
26	MW-2	Sample_Depth		0	0	0	0	0	0	0	0	7	7	100	7 p
27	MW-3	Temp	°C	0	0	0	0	0	0	0	0	7	7	100	7 p

Select

Parameters

Station

Export

Delete

Duplicate

New

Plot

Run

Close

The first 10 rows of the results table display a summary of the input options used for the trend analysis. These are defined under the **Options** tab. The rows below contain the results of the trend analysis. In the first section, it contains the station and parameter as well as the checks for data completeness. Following this section are the various tests depending on whether they are selected in the output options. The complete set of tests include:

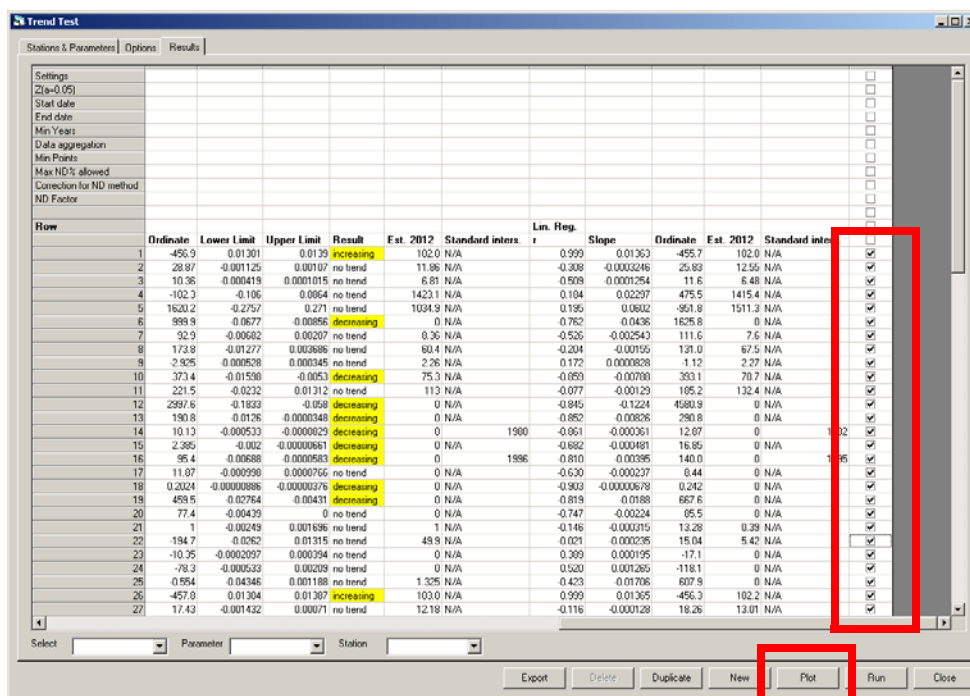
- Mann Kendall
- Seasonal Mann Kendall
- Sen's test
- Linear Regression
- Spearman's Rank Correlation

Significant increasing and decreasing trends are highlighted with a yellow background. If you would like to check the statistical test results, double click the respective row and you will be deferred to the Station & Parameter tab, where the station and parameter are automatically selected for you. Here you can select **Trend Results** from the **Show** combo box to view the statistical test results.

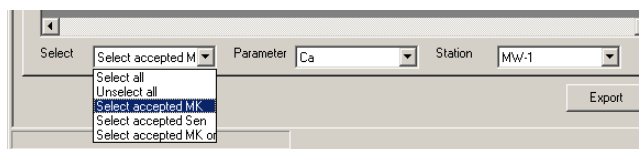
Creating Plots

Trend analysis plots can be quickly generated for each station and parameter combination in the results tab. In the results tab, scroll across to the end of the results

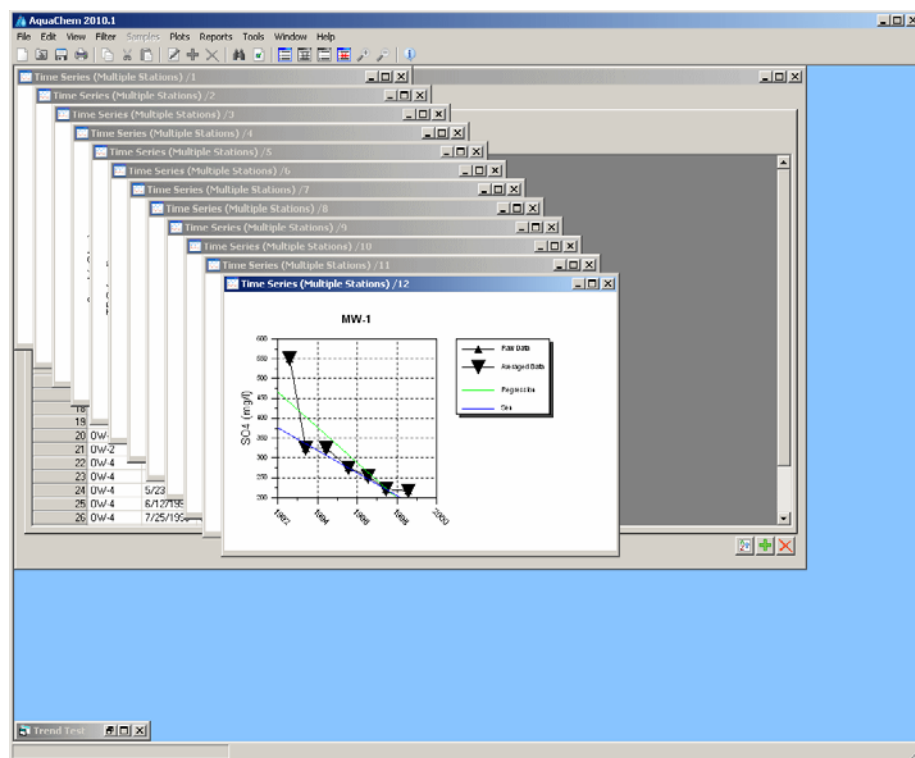
table, so that the last column is visible. This column will contain a selectable box in each cell. Select the desired rows to be plotted.



Note: The **Select** combo box, located at the bottom-right corner of the dialog, provides various options for automatically selecting multiple rows for plotting. For example, the **Select All** option will select all rows in the results table. The **Select accepted MK** option will only select those rows that meet the criteria for Mann Kendall analysis. Furthermore, you can filter the result list by parameter name (using the **Parameter** combo box) or station name (using the **Station** combo box)



Once the desired rows are selected, click the **[Plot]** button, located along the bottom of the **Trend Test** dialog. A time series plot displaying the linear regression and the Sen line will display for each selected row.



Saving the Results

To **Save** the trend analysis results, select the **Export** button , located at the bottom of the dialog box. You may chose from three file formats.

Comma Separated Values (*.CSV)	This format is useful when your table is very large (>65,000 rows) and you do not want to use the database format
Microsoft Excel (*.XLS)	This is the preferred format if you simply want to create a table for your report.
Microsoft Access (*.MDB)	This will create a new table in a specified database. This will allow you to query out specific results within the MS Access environment, e.g., if you want to know the percentage of insufficient data for a given parameter, the number of increasing versus decreasing trends etc.

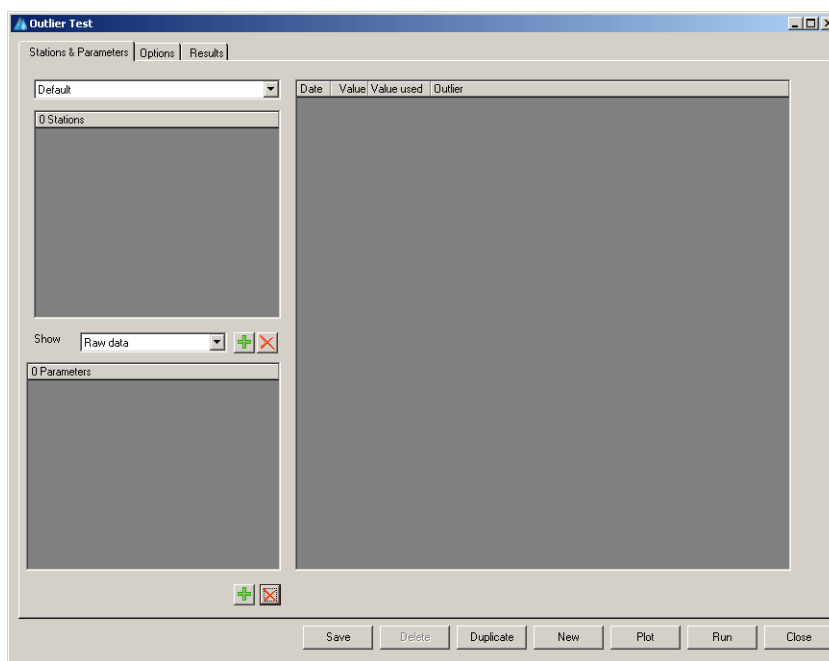
5.8.4 Outlier tests

Outliers are measurements that are extremely large or small relative to the rest of the data and, therefore, are suspected of misrepresenting the population from which they

were collected. Outliers may result from transcription errors, data-coding errors, or measurement system problems such as instrument breakdown. However, outliers may also represent true extreme values of a distribution (for instance, hot spots) and indicate more variability in the population than expected. Not removing true outliers and removing false outliers both lead to a distortion of estimates of population parameters. (Office of Environmental Information, U.S. Environmental Protection Agency, 2000).

AquaChem allows you to run outlier tests for multiple stations and parameters simultaneously.

Selecting this option from the **Statistics** sub-menu launches the following dialogue:



The workflow for performing batch outlier tests is similar to that of performing trend analysis test. Start by specifying the stations and which parameters to include in the outlier tests. This process is described on “Stations & Parameters” on page 280.

Exploring the Data

Before you run the actual outlier tests with potentially hundreds of stations and parameters, you should first explore the dataset and check whether there are stations or parameters which should be excluded from the analysis due to insufficient data.

Use the **Show** combo box, to display various information in the adjacent display area about the selected station and parameter

The show combo box provides the following options:

Raw Data	Shows all data points including analysis marked as outliers. Note that outliers will not appear in other previews, e.g., in plots
Statistics	Provides a short summary of statistics
Trend Test	Summarizes the outlier results for the current combination of parameters and stations. The information of each normality tests are shown: Dixon test, Discordance test, Rosner test, Walsh test. A description of thee tests together with directives for the calculations can be found in the <i>USE PA report QA/G-9 Guidance for Data Quality Assessment</i> .
Histogram Plot	Displays a time series plot for the selected parameter and station
Station Summary	Displays a summary of each station in the database that has an analyzed parameter value, including the Min, Max, Count, date of the First Sample and date o f the Last sample.
Parameter Summary	Displays a list of stations that contain data for the selected parameter, and associated information including the Min, Max, Count, date of the First Sample and date o f the Last sample.

Options

The options tab allows you to configure the various input and output options for the outlier test. Once you have specified the desired options, you can save the configuration by selecting the **[Save]** button from the row of buttons along the bottom of the dialog.

Outlier Test

Stations & Parameters Options Results

Input

Title: New Configuration

Description: Before you run the actual trend analysis, with potentially hundreds of stations and parameters, you should first explore the

ND Method: ND=MDL*Factor

ND Factor: 0.5

Max % ND: 10

Historic ND, ignore >: 10

Min Points: 5

Confidence (%): 95

Test for Normality (<=50 points): Studentized Range test

Test for Normality (>50 points): Studentized Range test

Confidence (% Normality test): 50

Output

Show Analyzed rows only: no

Show Percentile Lines: yes

Percentile: 95

Show Dixon's Test (3-25): yes

Show Discordance Test (<=50): yes

Show Rosner's Test (>=25): yes

Show Walsh's Test (>=60): yes

Symbol:

Configure the symbol used in the time series diagram

Save Delete Duplicate New Plot Run Close

Input

The input options allow you to configure various input parameters for the normality tests. Each option is described briefly below:

Title	Title of outlier test configuration
Description	A description of the outlier test configuration
ND Method	The method for handling non detect values
ND Factor	The factor by which all non detect concentrations are multiplied or constant value that is assigned to all non detect concentrations.
Max %ND	The maximum percent of non detects allowed in the dataset. If this threshold is exceeded, no test will be performed.
Historic ND, ignore	For long time intervals the detection limit method may have changed between early and recent data; more recent detection limits are often lower. When multiplying non detects by a factor, the data may become biased. Chose a ratio of historic/modern MDL above which you decide to ignore non detects. For example, if the value is set to 10 and the modern (minimum) MDL is 0.1 all non detects greater than <1 will be ignored.
Min Points	The minimum number of points required to run the outlier test. If this threshold is not reached, no test will be performed.
Confidence (%)	Confidence for all outlier tests

Test for Normality (<= 50 points)	Parametric outlier test requires that the underlying data is normally distributed. For each dataset, a normality test is run, and parametric tests are only run if the data is normally distributed. Chose the desired test for normality when the number of points is less than or equal to 50.
Test for Normality (>50 points)	Parametric outlier test requires that the underlying data is normally distributed. For each dataset, a normality test is run, and parametric tests are only run if the data is normally distributed. Chose the desired test for normality when the number of points is greater than 50.
Confidence (% , Normality test)	The confidence limit for the normality test

Output

The output options allow you to configure how the results of the outlier tests are displayed in the **Result** tab. The output options are described briefly below.

Show Analyzed Rows only	Choose whether or not to display rows where the data quality criteria have been matched. This is useful for tests of a large number of stations and parameters where many records do not reach the data quality criteria, e.g., there are not enough points or too many non detects.
Show Percentile Lines	Shows/hides control lines on time series plots
Percentile	Specify the percentile value that should be applied to the control line
Show Dixon's Test (3-25)	Show/hide the results of the Shapiro Wilk test in the result table
Show Discordance Test (<=50)	Show/hide the results of the Discordance Test in the result table.
Show Rosner's Test (>=60)	Show/hide the results of the Rosner's Test in the result table
Show Walsh's Test (>=60)	Show/hide the results of the Walsh's Test in the result table.
Symbol	Change the symbol of the data points displayed on outlier time series plots.

Results

Once you have specified the Stations & Parameters and have configured the input and out Options, you can run the outlier tests. To run the outlier tests, click the **[Run]** button located along the bottom of the **Outlier Test** dialog.

Depending on how many parameters and stations are include in the test, this may take a few minutes to complete. The results will display in the **Results** tab.

Row	Station	Parameter	Unit	Standard	min MDL	Non Detects	Outliers	MDL>0'S	Total samples	Valid samples	Years with Data	D
1	MW-1	Sample Depth		0	0	0	0	0	7	7	100	7 p
2	MW-1	Temp	°C	0	0	0	0	0	7	7	100	7 p
3	MW-1	pH_field		0	0	0	0	0	7	7	100	7 p
4	MW-1	Cond	uS/cm	0	0	0	0	0	7	7	100	7 p
5	MW-1	TDS	mg/l	0	0	0	0	0	7	7	100	7 p
6	MW-1	Ca	mg/l	0	10	1	14	0	7	7	100	7 p
7	MW-1	Mg	mg/l	0	0	0	0	0	7	7	100	7 p
8	MW-1	Na	mg/l	0	0	0	0	0	7	7	100	7 p
9	MW-1	K	mg/l	0	0	0	0	0	7	7	100	7 p
10	MW-1	Cl	mg/l	0	0	0	0	0	7	7	100	7 p
11	MW-1	HCO3	mg/l	0	0	0	0	0	7	7	100	7 p
12	MW-1	SO4	mg/l	0	0	0	0	0	7	7	100	7 p
13	MW-1	As_diss	ug/l	0	0.01	1	14	0	7	7	100	7 p
14	MW-1	Ba	mg/l	2	0	0	0	0	7	7	100	7 p
15	MW-1	Fe_diss	mg/l	0	0	0	0	0	5	5	100	5 p
16	MW-1	Hg_diss	ug/l	2	0.001	4	57	0	7	7	100	7 p
17	MW-1	Mn_diss	mg/l	0	0	0	0	0	4	4	100	4 p
18	MW-1	Pb_diss	ug/l	0	0.007	4	67	0	7	6	86	6 p
19	MW-1	Benzene	ug/l	0	1	2	29	0	7	7	100	7 p
20	MW-1	TCE	ug/l	0	0	0	0	0	7	7	100	7 p
21	MW-1	PCE	ug/l	0	2	4	57	0	7	7	100	7 p
22	MW-1	14C	% mod	0	0	0	0	0	5	5	100	5 p
23	MW-1	18O		0	0	0	0	0	7	7	100	7 p
24	MW-1	2H		0	0	0	0	0	7	7	100	7 p
25	MW-1	Tritium	T.U.	0	0	0	0	0	7	7	100	7 p
26	MW-3	Sample Depth		0	0	0	0	0	7	7	100	7 p
27	MW-3	Temp	°C	0	0	0	0	0	7	7	100	7 p

The first 9 rows of the results table display a summary the input options used for the outlier test. These options are defined under the **Options** tab. The rows below contain the results of the outlier tests. In the first section, it contains the station and parameter as well as the checks for data completeness. Following this section are the various tests depending on whether they are selected in the output options. The complete set of tests include:

- Dixon's Extreme Value Test
- Discordance Test
- Rosner's Test
- Walsh's Test

A description of these tests together with directives for the calculations can be found in the USEPA report QA/G-9> Guidance for Data Quality Assessment.

If you would like to check the statistical test results, double click the respective row and you will be deferred to the Station & Parameter tab, where the station and parameter are automatically selected for you. Here you can select **Outlier Results** from the **Show** combo box to view the statistical test results.

Creating Plots

Outlier test plots can be quickly generated for each station and parameter combination in the results tab. The procedure for creating plots for outlier tests is similar to creating plots for trend analysis tests. For more information, please refer to “Creating Plots” on page 287.

Saving the Results

To **Save** the outlier test results, select the **Export** button, located at the bottom of the dialog box. You may choose from three file formats. For more information on the available options, please refer to “Saving the Results” on page 289.

5.8.5 Tests for Normality

Many statistical tests and models are only appropriate for data that follow a certain distribution, in particular, the assumption of normality. There are many methods available for verifying the assumption of normality ranging from simple to complex. As recommended by the USEPA Guidance for Data Assessment document, there are five of the more common tests are built into AquaChem.

AquaChem allows you to perform normality tests for multiple stations and parameters simultaneously. The workflow for performing batch normality tests is similar to that of performing trend analysis test.

To perform normality tests, select **Reports > Statistics > Test for Normality**, from the AquaChem main menu. The **Test for Normality** dialog will appear on your screen, consisting of three tabs: Stations & Parameters, Options and Results.

Stations & Parameters

Start by specifying the stations and which parameters to include in the test for normality. This process is described on “Stations & Parameters” on page 280.

Exploring the Data

Before you run the actual test for normality, with potentially hundreds of stations and parameters, you should first explore the dataset and check whether there are stations or parameters which should be excluded from the analysis due to insufficient data.

Use the **Show** combo box, to display various information in the adjacent display area about the selected station and parameter

The show combo box provides the following options:

Raw Data	Shows all data points including samples marked as outliers. Note that outliers will not appear in other previews, e.g., in plots
-----------------	--

Statistics	Provides a short summary of statistics including the arithmetic mean, standard deviation, etc.
Trend Test	Summarizes the normality test results for the current combination of parameters and stations. The results of each normality tests is shown: Shapiro-Wil (W) test, Geary's test, D'Agostino's test.
Histogram Plot	Displays a histogram plot for the current station and parameter selection.
Station Summary	Displays a summary of each station in the database that has an analyzed parameter value, including the Min, Max, Count, date of the First Sample and date of the Last sample.
Parameter Summary	Displays a list of stations that contain data for the selected parameter, and associated data including the Min, Max, Count, date of the First Sample and date of the Last sample.

Options

The options tab allows you to configure the various input and output options for the normality test. Once you have specified the desired options, you can save the configuration by selecting the **[Save]** button from the row of buttons along the bottom of the dialog.

Test for Normality

Stations & Parameters | **Options** | Results

Input

Title	Default
Description	Default settings for normality tests
ND Method	ND=MDL*Factor
ND Factor	0.5
Max % ND	90
Historic ND, ignore >	100
Min Points	3
Confidence (%)	95
test log-normal	no

Output

Show Analyzed rows only	no
Geary's test (k=50)	yes
Studentized Range test (k=1000)	no
Shapiro-Wilk (W) test (n=50)	yes
Show Walsh's Test (b=80)	yes
Coefficient of variation	yes

Save Delete Duplicate New Plot Run Close

Input

The input options allow you to configure various input parameters for the normality tests. Each option is described briefly below:

Title	The title of the trend test configuration
Description	A description of the trend test configuration
ND Method	Method for handling non detect values
ND Factor	The factor by which all non detect concentrations are multiplied or the constant value that is assigned to all non detect concentrations.
MAx % ND	The maximum percent of non detects allowed in the dataset. If this threshold is exceeded, no test will be performed.
Historic ND, ignore >	For long time intervals, the method detection limit may have changed between early and recent data, and more recent detection limits are often lower. When multiplying non detects by a factor, the dataset may become biased. Chose a ratio of historic/modern MDL above which you decide to ignore non detects. For example, if the value is set to 10 and the modern (minimum) MDL is 0.1, all non detects greater than < 1 will be ignored.
Min Points	The minimum number of points. If this threshold is not reached, the test will not be performed.
Confidence (%)	The confidence for normality tests.
test log-normal	Select whether you want to run the test on lognormally transformed data.

Output

The output options allow you to configure how the results of the normality test are displayed in the Result tab. The output options are described briefly below.

Show Analyzed Rows only	Choose whether or not to display rows where the data quality criteria has not been matched. This is useful for tests of a large number of stations and parameters where many records do not reach the data quality criteria, e.g., there are not enough points to too many non detects.
Geary's test (≤ 50)	Show/hide Geary's normality test results
Studentized Range test (≤ 1000)	Show/hide Studentized Range test results
Shapiro-Wilk (W) test (≥ 50)	Show/hide Shapiro Wilk (W) test results

Show Walsh's Test (>=60)	Show/hide Walsh's test results
Coefficient of variation	Show/hide the results of the calculated coefficient of variation

Results

Once you have specified the Stations & Parameters and have configured the input and output Options, you can run the normality test. To run the normality test, click the **[Run]** button located along the bottom of the **Test for Normality** dialog.

Depending on how many parameters and stations are include in the test, this may take a few minutes to complete. The results will display in the **Results** tab.

The first 7 rows of the results table display a summary the input options used for the normality test. These options are defined under the **Options** tab. The rows below contain the results of the normality tests. In the first section, it contains the station and parameter as well as the checks for data completeness. Following this section are the various tests depending on whether they are selected in the output options. The complete set of tests include:

- Geary's test
- Studentized Range test (t-test)
- Shapiro-Wilk test (w-test)
- d'Agostino's test, and
- Coefficient of Variation

A description of these tests together with directives for the calculations can be found in the EPA report QA/G-9 Guidance for Data Quality Assessment, sections 4.2 and following.

The result of each test may be **True**, **False** or **<N**. A true result indicates that the dataset can be considered to have a normal distribution according to the respective test. A false result indicates that the data cannot be considered to have a normal distribution. A **<N** result, where N is the minimum points required, means that there is not sufficient data points to perform the normality test.

If you would like to check the statistical test results, double click the respective row and you will be deferred to the Station & Parameter tab, where the station and parameter are automatically selected for you. Here you can select **Trend Results** from the **Show** combo box to view the statistical test results.

Creating Plots

Histogram plots can be quickly generated for each station and parameter combination in the results tab. The process for creating plots for normality tests is similar to that for trend analysis. For more information, please refer to "Creating Plots" on page 287.

Saving the Results

To **Save** the trend analysis results, select the **Export** button, located at the bottom of the dialog box. For more information on the available options, please refer to “Saving the Results” on page 289.

5.8.6 Alert Levels

An **Alert Level (AL)** refers to an upper tolerance level for a constituent which can be used as an early warning indicator of a potential violation of a water quality standard. In AquaChem, Alert Levels can be calculated for one or more constituents using baseline monitoring data. The alert levels may then be compared with future detection monitoring data to determine statistically significant increases of constituents in groundwater.

AquaChem supports three methods for calculating statistically significant alert levels using historical data: **Probability**, **STD x Factor** and **Tolerance Interval**. Each method must satisfy the following criteria before it can be deemed appropriate for calculating alert levels:

- The historical dataset must pass a normality test
- The percentage of non detects in the dataset cannot exceed a specified maximum percentage.
- The dataset must equal or exceed a specified number of data points.

If the dataset does not meet one of the above requirements, AquaChem allows you to determine alert levels using alternative methods, *e.g.*, MDL x Factor, Max value.

To calculate **Alert Levels**, select **Reports > Statistics > Alert Levels** from the main menu.

The **Alert Levels** dialog will appear on your screen consisting of three tabs: **Stations & Parameters**, **Options** and **Results**.

Stations & Parameters

The workflow for selecting stations and parameters for the alert level calculations is similar to that of selecting stations and parameters for trend analysis tests. For more information, please see “Stations & Parameters” on page 280.

Exploring the Data

Before you run batch calculations with potentially hundreds of stations and parameters, you should first explore the dataset and check whether there are stations or parameters which should be excluded from the analysis due to insufficient data.

Use the **Show** combo box, to display various information in the adjacent display area about the selected station and parameter

The show combo box provides the following options:

Raw Data	Shows all data points including analysis marked as outliers. Note that outliers will not appear in other previews, e.g., in plots
Statistics	Provides a short summary of statistics including the arithmetic mean, standard deviation, etc.
Alert Level values	Displays the calculated alert level for the current station and parameter selection. The calculation is based on the input options specified under the Options tab. For more information on the options for Alert Levels, please refer to “Options” on page 300.
Time Series Plot	Displays a time series plot for the current station and parameter selection. The upper tolerance level is also represented graphically on the time series plot.
Station Summary	Displays a summary of each station in the database that has an analyzed parameter value, including the Min, Max, Count, date of the First Sample and date of the Last sample.
Parameter Summary	Displays a list of stations that contain data for the selected parameter.

Options

The options tab allows you to configure the various input and output options for calculating alert levels. Once you have specified the desired options, you can save the configuration by selecting the **[Save]** button from the row of buttons along the bottom of the dialog.

Input

The input options allow you to configure various input parameters for the alert level calculations. Each option is described briefly below:

Title	The title of the alert level configuration
Description	A description of the alert level configuration
Alert Level method	<p>There are three methods for calculating alert levels: Probability, STD x Factor, Tolerance Interval.</p> <p>Probability: The alert level is calculated as the concentration where the percentage of exceeding this value is lower than a specified confidence limit, e.g., 1%. When this method is used, you must specify the desired probability percentage in the field below.</p> <p>STD x Factor: The alert level is calculated as the mean plus the standard deviation multiplied by a user specified factor. When this method is used, you must specify the desired factor in the field below.</p> <p>Tolerance Interval: The alert level is calculated as the upper tolerance limit. The tolerance interval includes the 95% of the data with the specified confidence. When this option is selected, a Confidence field will appear where you can specify the desired confidence value. For more information on the tolerance interval method, please refer to “Appendix D: Calculating Alert Levels” on page 425.</p>

Test for	Calculate one sided (Upper AL) or two sided (Upper and Lower AL) alert levels. Parameters which have a lower and upper water standard, such as pH or alkalinity, should be calculated as two sided. For parameters where one is concerned about high values, the one sided alert level is sufficient.
Alternative AL if ND>90%	<p>If the maximum percentage of non detects in the dataset is exceeded, alert levels will not be calculated using the specified alert level method. In this case, alert levels may be determined using an alternative method. If MDL option is selected, the alert level will be calculated as the product of the method detection limit and a user specified factor. If the Max value option is selected, the alert level will be calculated as the product of the maximum value in the dataset and a user specified value.</p> <p>The factors for each options may be specified in the field below, labeled Factor (ND% exceeded).</p>
Alternative AL if data is not normal	<p>If the dataset is not of normal distribution, alert levels will not be calculated using the specified alert level method. Instead, alert levels may be determined using an alternative method. The Regular AL option allows you to override this requirement and calculate the alert levels using the specified method despite it not being of normal distribution. If the MDL option is selected, the alert level will be calculated as the product of the method detect limit and a user specified factor. If the Max value option is selected, the alert level will be calculated as the product of the maximum value in the dataset and a user specified value.</p> <p>The factors for the MDL and Max Value option may be specified in the field below, labeled Factor (not normal).</p>
Start Date	Specify the starting date from which the alert level is to be calculated.
End Date	Specify the end date from which the alert level is to be calculated.
ND Method	Specify the method for handling non detects. The MDL x Factor option allows you to use calculate concentrations by multiplying the method detection limit by a user specified factor. The Const option allows you to assign a constant value to all non detect measurements. The factor or constant value can be specified in the field below.
Max % ND	Specify the acceptable maximum percentage of non detects in the dataset from which the alert levels will be calculated. If the percentage of non detects exceeds this value, the alert levels will not be calculated using the specified alert level method.
Correction for 15% ND% <50%	Specify the correction method for calculating the mean and standard deviation of the dataset, if the percentage of non detects is greater than 15% and <50%.

Correction for ND%>50%	Specify the correction method for calculating the mean and standard deviation if the percentage of non detects is greater than 50%
Historic ND, ignore >	Data sets with large time intervals may show a decrease in the method detection limit over time. Data sets that exhibit this trend may become biased if non detects are multiplied by a factor. To handle this scenario, you can specify a ratio of historic/modern MDL above which you decide to ignore non detects. For example, if the value is set to 10, and the modern (minimum) MDL is 0.1, all non detects greater than <1 will be ignored.
Min Points	Specify the minimum number of data points required for alert level calculations. If the specified threshold is not satisfied, the alert level will not be calculated.
Max Points	Specify the maximum number of points to be included in the alert level calculation. For example, if this value is set to 8 and there are more samples than the maximum number of points in the dataset, only the 8 most recent samples will be used to calculate the alert level.
Test for normality (<=50 points)	Specify the desired test for normality if the dataset contains less than or equal to 50 points. You may choose from Studentized Range test, Geary's test or the Shapiro-Wilk test. For more information on normality tests, please refer to "Tests for Normality" on page 295.
Test for normality (>50 points)	Specify the desired test for normality if the dataset consists of more than 50 points. You may choose from Studentized Range test, Geary's test or the Shapiro-Wilk test. For more information on normality tests, please refer to "Tests for Normality" on page 295.
Confidence (% , normality)	Specify the confidence value for the specified normality test.

Output

The output options allow you to configure how the results of the alert level calculations are displayed in the alert levels plots. The output options are described briefly below.

Standard	Choose which standard to show on the control plot.
Show standard line	Show/hide water quality standard line in control plot
Show alert levels	Show/hide alert levels as lines in control plot
Symbol	Specify the desired symbol for showing alert level data points on the control plot.

Results

Once you have specified the Stations & Parameters and have configured the input and output Options, you can run the alert level calculations. To run the alert level calculations, click the **[Run]** button located along the bottom of the **Alert Level** dialog.

Depending on how many parameters and stations are included in the test, this may take a few minutes to complete. The results will display in the **Results** tab.

The first rows of the results table display a summary of the input options used for the alert level calculations. These options are defined under the **Options** tab. The rows below contain the details of the alert level calculations. In the first section, it contains the station and parameter as well as the checks for data completeness and normality. Following this section are the mean, standard deviation and calculated alert levels.

If you would like to check the calculations in more detail, double click the respective row and you will be deferred to the Station & Parameter tab, where the station and parameter are automatically selected for you. Here you can select **Alert Level Values** from the **Show** combo box to view the specifics of the alert level calculation.

Plotting Alert Levels

Once the alert levels have been calculated, you can display the upper (and lower) alert level limit with the corresponding background dataset on a time series plot. Plots can be generated from the Result tab.

Under the **Plot** column, in the **Results** tab, select the desired rows to be plotted (see image below)

Note: The **Select** combo box, located at the bottom-right corner of the dialog, provides various options for automatically selecting multiple rows for plotting. For example, the **Select All** option will select all rows in the results table. You can filter the result list by parameter name (using the **Parameter** combo box) or station name (using the **Station** combo box)

To create the alert level plots, click the **[Plot]** button, located at the bottom of the dialog. The series of plots will display on your screen. From here you can print the plots or save the plot configuration for viewing at a later time.

Alert Levels

Stations & Parameters | Options | Results

Settings

Method

Start date baseline

End date baseline

Min Points

Max Points

Max ND % allowed

Correction for ND method

ND Factor

Normality test n<=50

Normality test n>50

Row	%	Total samples	Valid samples #	%	Data checks	Normality test	Correction	Mean	STD	Upper AL	Plot
1	N/A	5	5	100	passed	False		45.2	2.55	49.4	<input type="checkbox"/>
2	N/A	5	5	100	passed	False		8.98	0.3834	9.61	<input checked="" type="checkbox"/>
3	N/A	5	5	100	passed	False		24.9	2.153	28.46	<input checked="" type="checkbox"/>
4	N/A	5	5	100	passed	False		5.66	0.568	6.6	<input type="checkbox"/>
5	N/A	5	5	100	passed	True		9.54	1.606	12.18	<input checked="" type="checkbox"/>
6	N/A	5	5	100	passed	False		7.38	0.439	8.1	<input checked="" type="checkbox"/>
7	62	5	5	100	passed	True		284	300.5	778.2	<input checked="" type="checkbox"/>
8	31	5	5	100	passed	True	Cohen's	13	9.17	28.1	<input checked="" type="checkbox"/>
9	85	3	3	100	passed	True		9.35	8.05	22.6	<input type="checkbox"/>

Select Parameter Station

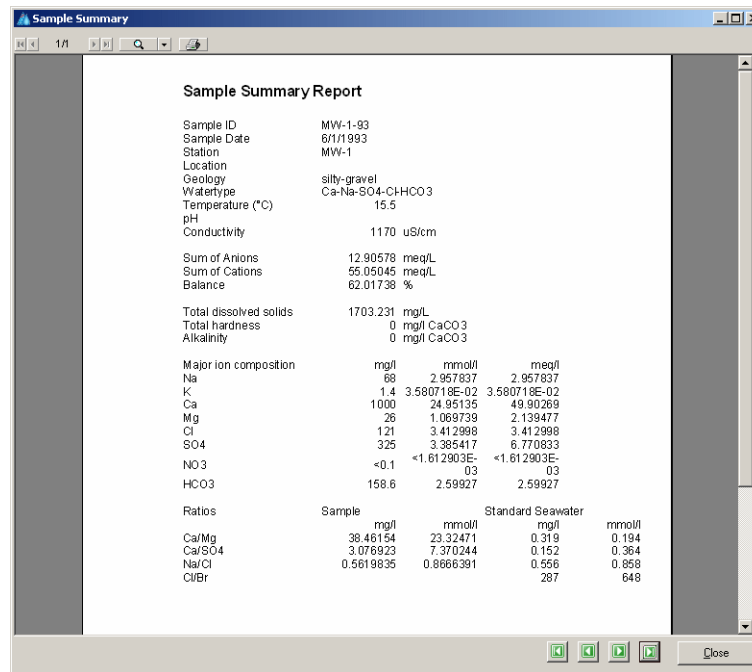
Export Delete Duplicate New Plot Run Close

5.9 Sample Summary

The **Sample Summary** report is formatted differently from the other reports described up to this point. This report was designed using the **Report Designer**; as such, you have the option of modifying the appearance of this report, including the layout, available fields, and printing template. For more details on how to modify this report template, please see the **Report Designer** section at the end of this chapter.

The **Sample Summary** report provides a general overview of a single sample including major ions, hydrochemical facies (e.g. Na-Cl), calculated hardness, ion balance, ion ratios, etc. The parameter values are read directly from the **Sample Details** window for each sample; this report shows both measured and calculated values.

When you select the **Reports** from the main menu and then **Sample Summary**, the following **Sample Summary** report window will appear.



The report window for user-defined reports (as shown in the figure above) has a few more options available at the top and bottom of the report window:



The **Scroll arrows** are only available when you have a report that contains multiple pages. The page count shows the current page number, out of the total number of pages available.

The **Zoom** (magnifying) button allows you to change the zoom size of the window. When you press the down arrow beside the magnifying glass, you will see several window size options. Simply choose the desired zoom size and the preview window should be automatically refreshed.

The **Print** button sends the reports for all selected samples to the printer.



The **Show fields** option is only available if the Report is linked to a printing template (done using the **Report Designer**). To enable this option, place a check mark in this box. You will then see the following dialogue appear in the middle of your display, to the left of the preview window:

In these fields, you can enter the appropriate descriptive information that should appear in the template. As the information is entered, the report preview window should be automatically updated.

Parameter	Value
Description	Project Description.....
DATE	Report Date.....
CLIENT	Client Info.....
PROJECT NO	Project Number Info.....

☒ Show fields

The **Scroll arrows** on the bottom of the report window are only available when only one sample is selected. They allow you to scroll through the entire dataset.

NOTE: You may generate single-page and multi-page reports. Selecting multiple samples and generating a **Sample Summary** report will generate an individual report for each selected sample, allowing you to scroll through them using the scroll buttons at the TOP of the report window. Printing the report at this stage will print a report for ALL selected samples. Alternately, if you generate the **Sample Summary** report while only one sample is selected, the report will be created for that samples and you will be able to scroll through all samples in the project using the green scroll buttons at the BOTTOM of the report window. Printing the report will produce a printout of this sample ONLY.

Clicking the **[Close]** button closes the report.

This report will be edited in **Report Designer**.

NOTE: When you save user-defined reports, the report template information will not be saved; only the report contents are saved to a file.

5.10 Thermometers

This report lists the geochemical thermometers and their estimated temperatures which are calculated using the formulas defined in the **Thermometers** tab of the **Database**. This report may be edited in **Report Designer**.

5.11 Isotopes

This report lists the concentrations of isotopes ^{18}O and ^2H as well as infiltration temperature and height for ^{18}O which are calculated according to formulas specified in **Calculations** tab of the **Database**. This report may be edited in **Report Designer**.

5.12 Report Designer

As mentioned earlier in this chapter, AquaChem allows you to create two types of reports:

- The pre-defined (hardwired) reports, and
- The user-defined reports

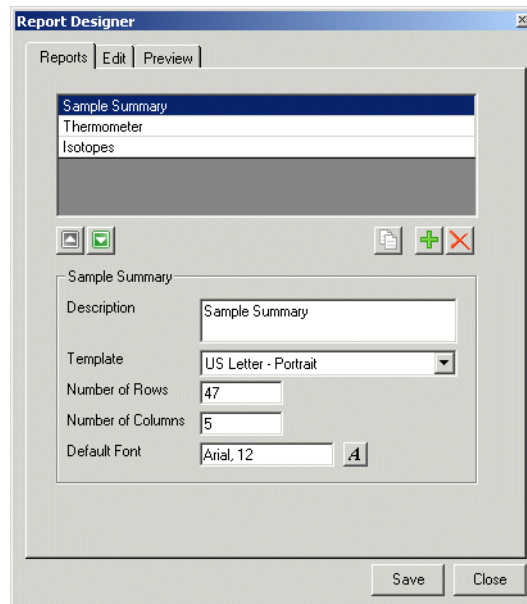
The hardwired reports cannot be modified, nor can their layout be changed. These reports include: **Compare Sample**, **Mix Samples**, **Water Quality Standards**, **Reliability Check**, **Compare Duplicates**, **Rock Source Deduction**, **Summary Statistics**, and **Correlation Matrix**.

The user-defined reports (ex. **Sample Summary**) can be modified using the **Report Designer**.

Using the **Report Designer** you may design and customize up to 10 new reports for your own project needs. The report may include any combination of database parameters and function (calculation) results. Once a new report has been created, it can be selected from the **Reports** menu, below the **Isotopes** report.

5.12.1 General Features

When you select **Report Designer** from the **Reports** menu, the following dialogue will appear. The **Report Designer** dialogue is separated into three tabs: **Reports**, **Edit**, and **Preview**.



Reports

Provides a list of the reports available in the current database template, and general information on each report (Description, Selected Printing template, report size, and selected font).

Edit

Provides options for designing the report layout.

Preview

This tab provides you with a preview of the selected report.

You will also find the following buttons in the **Report Designer** dialogue.



The up-arrow button allows you to change the order of the selected reports upwards as they appear in the **Reports** menu (in the main menu).



The down-arrow button allows you to change the order of the selected reports downwards as they appear in the **Reports** menu (in the main menu).



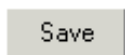
The **Duplicate** button clones the selected report. This allows you to quickly modify an existing report, without having to create a new one from scratch.



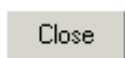
The **Add** button creates a new report.



The **Delete** button deletes the selected report.



The **[Save]** button at the bottom of the dialogue saves the current report information.



The **[Close]** button closes the Report Designer dialogue.

In order to quickly learn the functionality of the **Report Designer**, an example has been provided below.


5.12.2 Designing a New Report - Example

Before customizing a report you should create a spreadsheet-style layout of this report using a program such as Excel. Decide what fields you want to use, their placement, and how wide the cells need to be. This will speed-up the process when using the Report Designer in AquaChem. Below is the layout that will be used for this simple example report.

	A	B	C	D	E
1	General Report				
2					
3	Sample ID				
4	Sampling Date				
5	Location				
6					
7	Major Ions				
8		mg/L		mg/L	
9	Ca		CL		
10	Mg		SO4		
11	Na		HCO3		
12					
13	Sum Anions		Sum Cations		
14	Ion Balance				
15					
16	Ratio	mg/L			
17					
18	NA/CL				

To create a new report, follow the steps as described below.

If you have not already done so, load the **Report Designer** from the **Reports** menu. Ensure you are viewing the **Reports** tab (the first tab).

➡  button to create a new report. A new line will be added below the **Isotopes** Report, with the name: **#New Report**.

☞☞ **#NewReport** and enter a name for this new report.

type: **Example**

☞ **<Enter>** (on your keyboard)

In the lower half of this dialogue:

type: **An Example Report** (in the **Description** field)

☞ **'none'** for the Template

type: **20** (the **Number of Rows** for this report)

type: **5** (the **Number of Columns** for this report)

☞  button to change the **Default Font** for this report

☞ Times New Roman / Regular / 10 font size

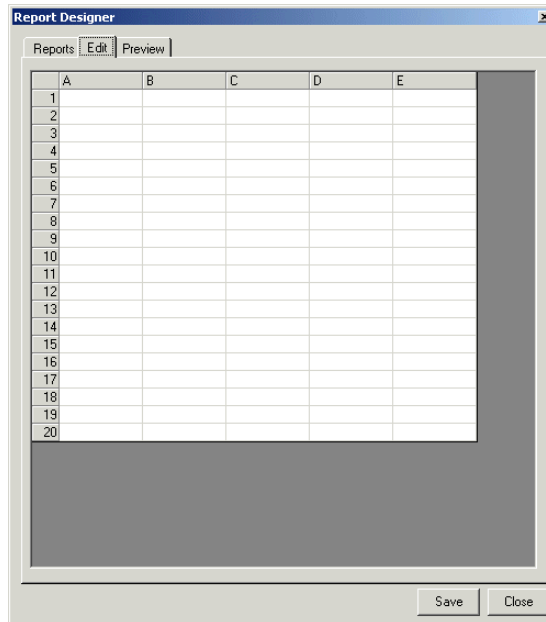
☞ **[OK]**

☞ **[Save]** (in the lower right corner of the dialogue) to save this report.

You can now proceed to add fields to this report.

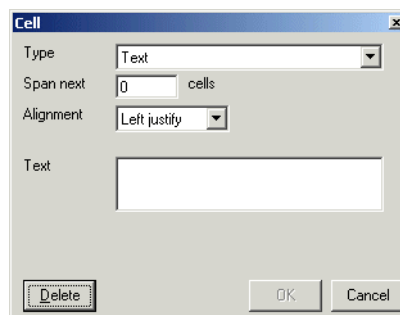
☞ **Edit** tab (from the top of the dialogue).

This will present you with a blank grid (spreadsheet), containing 20 rows and 5 columns as shown below.




You will see that the rows are numbered and the columns are lettered (similar to MSEXcel) to allow you to quickly locate cell locations in the grid. At this point you may begin to add values to the cells. You may fill cells with text, database parameters, or with a function (calculation).

☞ **Double-click** on the cell in the upper-left corner (Cell **A1**) and the **Cell** dialogue (as shown below) will appear:



This dialogue allows you to control the contents of the selected cell. Cell contents must be assigned one cell at a time. Below are the options available for each cell:

Type

Click on the  button and you may choose from the following cell types:

Text: Enter the desired text for the cell

Sample Description: choose from a list of Sample Description parameters

Station Description: choose from a list of Station Description parameters

Measured/Modeled Value: choose from a list of Measured or Modeled parameter values.

Ratio: choose two parameters to show as a ratio

Guideline Level1: displays the current guideline level 1 for the specified parameter

Guideline Level2: displays the current guideline level 2 for the specified parameter

Guideline Level3: displays the current guideline level 3 for the specified parameter

Function Value: choose from a list of available calculations

Range Name: select from a list of available Ranges

Thermometers: select from a list of Geothermometers

Span next

This option is useful for headings and titles, where the text may not fit into one cell. If the cell to the right of the selected cell is empty, you may span several cells into one. Simply enter a cell number in this field (ex. Span 2 cells).

Alignment

Choose from Left, Center, or Right-Justified.

The three fields described above are common to all data types listed. Below, all data types are listed along with the descriptions of the other fields that go with them.

Text

Text

Enter the text you wish to appear in this cell. Used for captions that do not change (e.g. "Station ID")

Sample Description

Sample Parameter

Use the combo box to select a sample parameter. The value of this cell will change depending on what sample is loaded into the report.

Station Description

Station Parameter

Use the combo box to select a station parameter. The value of this cell will change as you move between samples that are assigned to different stations.

Measured/Modeled Parameter

Parameter

Use the combo box to select a measured or a modeled parameter. The cell will reflect the value of this parameter for the sample that is loaded into the report.

Unit

Use the combo box to select a unit for the parameter (becomes inactive if the chosen parameter is unitless - e.g. pH).

Format

Use the combo box to select the numeric format for the parameter value.

Ratio

Numerator

Use the combo box to select the parameter to serve as the numerator of the ratio.

Denominator

Use the combo box to select the parameter to serve as the denominator of the ratio.

Units

Use the combo box to select the units for the parameters used as the numerator and the denominator. The ratio may change depending on the chosen units (e.g. the ratio of the parameter values in g/l will be different from the ratio of the parameter values in mol/l).

Format

Use the combo box to select the numeric format for the ratio.

Guideline Level 1, 2, and 3

Guideline Level

Use the combo box to select the parameter for which the selected guideline level will be displayed. The value of the cell will stay the same regardless of the sample loaded into the report.

Format

Use the combo box to select the format for the cell value.

Function Value

Function

Use the combo box to select a function, the result of which will be displayed in the cell. The functions are defined in the **Calculations** tab of the **Database**.

Unit

Use the combo box to select the unit for the parameters used in the function. The result of the calculations may be different depending on the units used.

Format

Use the combo box to select the format for the result value.

Range Name

Parameter

The combo box lists the parameters that have the range(s) defined. The cell will display the name of the range that contains the value of the specified parameter for the current sample (e.g. Natural aquifer range).

Thermometer

Thermometer

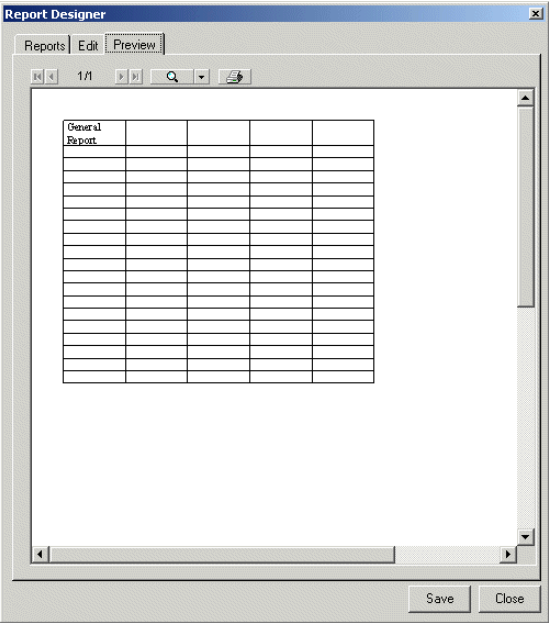
Use the combo box to select the thermometer function. The thermometer calculations are defined in the **Thermometers** tab of the **Database**.

“Designing a New Report” continued

You will now enter a Report Title for cell A1:

- ☞ **Text** (from the list beside the **Type** field)
- type: **1** in the **Span next ... cells** field
- ☞ **Left justify**, for the cell alignment
- type: **General Report** (beside the **Text** field)
- ☞ **[OK]**

You should now see an entry under the **Edit** tab for Cell **A1**. You may switch to the **Preview** tab to see a preview of how the report will appear.



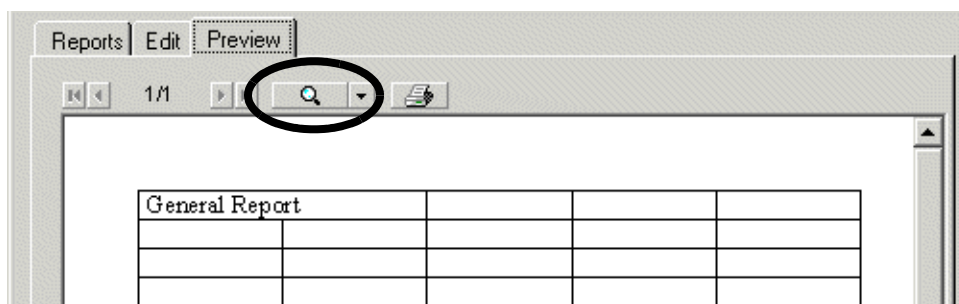
As you can see, the words take up two lines in the cell. To have the words fit on one line you can span this cell into the adjacent one.

Switch back to the **Edit** tab

☞☞ **Double-click** on the cell **A1** to produce the **Cell** dialogue

type: **2** in the **Span the next ... cells**

Switch to the Preview tab again. Now it should look similar the picture below (use the magnifying glass menu to adjust the view size):



The text colors of the cells in the **Edit** dialogue are representative of the cell types. The following summary provides a list of what each cell color indicates of the cell type:

Black: Text cells

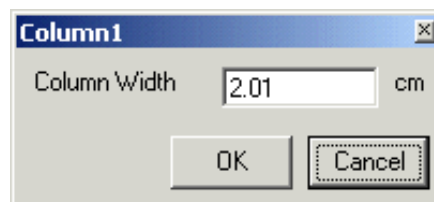
Green: Sample and Station Description parameters

Blue: Measured and Modeled parameters, and Parameter Ratios

Grey: Function (Calculated) values, Thermometers, and Ranges

Red: Guideline Levels

If you want to change the column widths, simply drag the column to the desired width in the spreadsheet. Alternatively, you can select the column then right-click the highlighted column and select **Set Column Width**. Enter the precise value of the column width in the pop-up dialogue and press **[OK]**. Then load the **Preview** window to see the refreshed report.



The font for the report may be customized in two places:

In the **Reports** tab, the **Default Font** for the entire report is defined; all items on the report will use this font, size and style.

A default font may also be assigned on a per row basis. To do so, select an entire row and right-mouse click on it and select **Font**. Select the desired **Font** from the dialogue that appears, and click **[OK]**. This font will be applied to an entire row. This is practical for assigning fonts to column headings in the report.

You can also insert and delete any selected row, column, or cell by highlighting the desired element, right-clicking on it, and choosing the appropriate action.

You will now proceed to define the remaining cells in the report, as per the pre-designed layout:

☞☞ Cell **A3**

☞ Select **Text** from **Type** field

type: **Sample ID**

☞ **[OK]**

☞☞ Cell **A4**

☞ Select **Text** from **Type** field

type: **Sampling Date**

☞ **[OK]**

☞☞ Cell **A5**

☞ Select **Text** from **Type** field

type: **Location**

☞ **[OK]**

Using the same procedure, fill in the remaining text fields as per the layout above, in the appropriate cell locations:

<u>Cell</u>	<u>Text Entry</u>
A7	Major Ions
A9	Ca
A10	Mg

A11	Na
A13	Sum Anions
A14	Ion Balance
A16	Ratio
A18	Na/Cl
B8	mg/l
B16	mg/l
C9	Cl
C10	SO4
C11	HCO3
C13	Sum Cations
D8	mg/l

Click on the **Preview** tab to see if any caption is too big for its cell. Go back to the **Edit** tab and if there were such captions, resize the column they are in so that they only take up one line. Do not worry if some parameter names (such as “SAMPLE_DATE”) take up a lot of space - gauge the length of the value.

Once you are finished you should save the changes that have been made so far.

- ☞ **[Save]** in the lower-right corner of the **Report Designer** dialogue, to save the report design changes up to this point.

You can now add in the parameter value cells and function values.

☞☞ **Cell B3**

- ☞ **Sample Description** from the combo box beside cell **Type**
- ☞ **SAMPLEID** from the combo box beside **Sample Parameter** field
- ☞ **[OK]**

☞☞ **Cell B4**

- ☞ **Sample Description** from the combo box beside cell **Type**
- ☞ Select **SAMPLE_DATE** from the combo box beside **Sample Parameter** field

☞ [OK]

☞☞ Cell B5

☞ **Station Description** from the combo box beside cell **Type**

☞ Select **LOCATION** from the combo box beside **Station Parameter**

☞ [OK]

Notice the color of the cells under the **Edit** tab in the **Report Designer** dialogue for these new cells; green cells indicate that these cells contain **Station/Sample Description** parameters.

☞☞ Cell B9

☞ **Measured /Modeled Value** from the combo box beside cell **Type**

☞ **Ca** from the combo box beside **Parameter**

☞ **mg/l** from the **Unit** field

☞ [OK]

☞☞ Cell B10

☞ **Measured Modeled Value** from the combo box beside cell **Type**

☞ **Mg** from the combo box beside **Parameter**

☞ **mg/l** from the **Unit** field

☞ [OK]

☞☞ Cell B11

☞ **Measured Modeled Value** from the combo box beside cell **Type**

☞ **Na** from the combo box beside **Parameter**

☞ **mg/l** from the **Unit** field

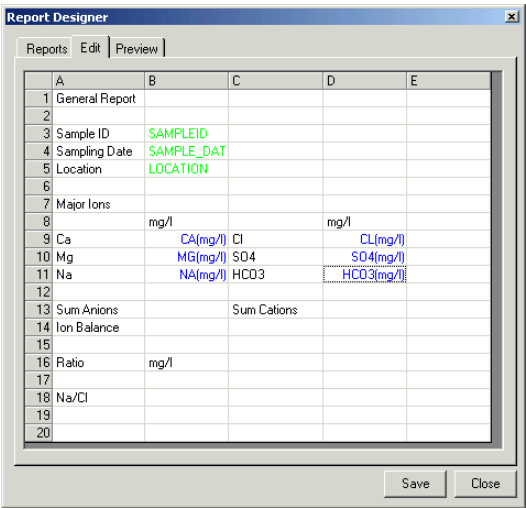
☞ [OK]

Notice the color of the cells in the **Edit** dialogue; blue cells indicate that these cells contain **Measured values**.

Repeat this step for the remaining ions by choosing the appropriate parameter and placing this in the cell immediately beside the text cell for that parameter.

<u>Cell</u>	<u>Measured Value</u>
D9	Cl
D10	SO4
D11	HCO3

Once you have finished this step, your display should be similar to the one shown below:



☞ **[Save]** in the lower-right corner of the **Report Designer** dialogue to save the report design changes up to this point.

You will now add function values for the **Sum Anions**, **Sum Cations**, and **Ion Balance**

- ☞☞ Cell **B13**
- ☞ **Function Value** from the combo box beside cell **Type**
- ☞ **Left Justify** from the **Alignment** field

☞ Select **Sum of Anions** from the combo box beside **Function**

☞ [OK]

☞☞ Cell **D13**

☞ **Function Value** from the combo box beside cell **Type**

☞ **Left Justify** from the **Alignment** field

☞ Select **Sum of Cations** from the combo box beside **Function**

☞ [OK]

☞☞ Cell **B14**

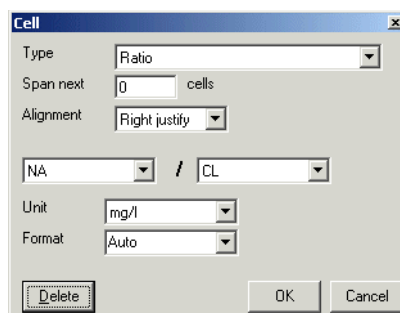
☞ **Function Value** from the combo box beside cell **Type**

☞ **Left Justify** from the **Alignment** field

☞ **E. N.** (Electroneutrality) from the combo box beside **Function**

☞ [OK]

Notice the color of the cells in the **Edit** dialogue; grey cells indicate that these cells contain **Function (Calculated) Values**. Finally, you will add a **Ratio** to Cell B18 showing a parameter ratio of Na / Cl.



☞☞ Cell **B18**

☞ Select **Ratio**, from the combo box beside cell **Type**

☞ **Na** for the first parameter

☞ **Cl** for the second (right) parameter

☞ **mg/l** for the units

☞ [OK]

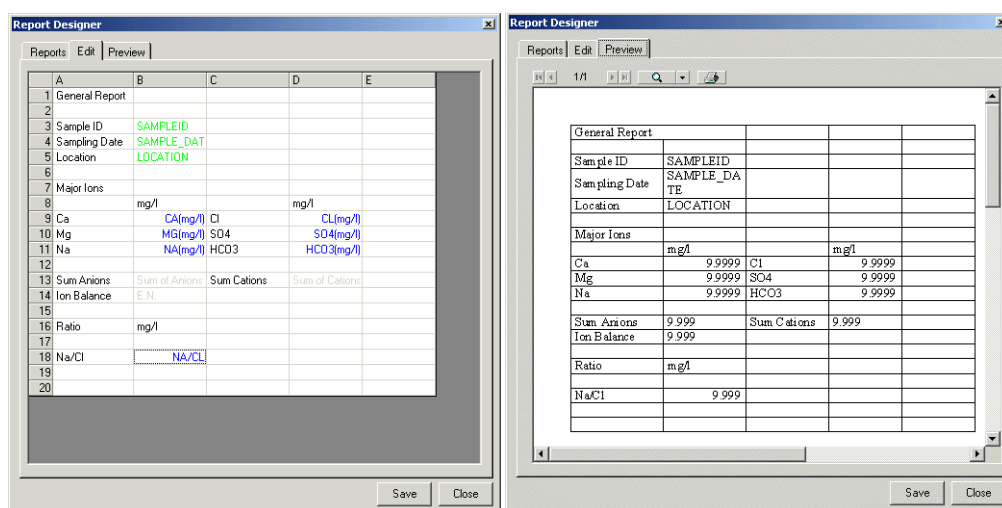
Notice the color of the cells in the **Edit** dialogue; blue cells indicate that this cell contains a **Parameter Ratio**.

☞ **[Save]** in the lower-right corner of the report designer dialogue to save all changes to this report.

Once you have designed the report, click on the **Preview** tab to see a page preview.

Edit Tab

Preview Tab



This concludes the steps required to design a report. Feel free to add new cell values, functions, parameters, etc. in order to see the full capabilities of the **Report Designer**.

Once you are finished in the **Report Designer**, press **[Close]** to return to the main window of AquaChem. The newly created “**Example Report**” will now be available under **Reports** in the AquaChem main menu.

6

Tools

The **Tools** menu provides access to the AquaChem data analysis tools, including Converters, Calculators, LookUp Tables, and access to the PHREEQC modeling interface.

The following **Calculators and Converters** are available:

- Aggregate Samples
- AquaChem Function
- Calculate facies
- Corrosion & Scaling
- Decay Calculator
- Find Missing Major Ion
- Formula Weight Calculator
- Oxygen Solubility
- Volume Concentration Converter
- Retardation
- Special Conversions
- Species Converter
- Unit Calculator
- UTM Conversion

The following **QA/QC** checks are available:

- Compare Duplicates
- Control Intervals
- Highlight Duplicates
- Highlight Outliers
- Holding Times
- Manage Duplicates
- Method Detection Limit
- Reliability Check
- Total/Dissolved Comparison

The following **LookUp Tables** are available:

- Degradation Rates
- PHREEQC Phases
- Periodic Table
- Water Standards
- Time Series
- Browse Database

Finally, under the **Modeling** options you may:

- Calculate Saturation Indices and Activities
- Calculate pH
- Calculate Eh
- Alk > HCO₃, CO₃
- Equilibrate with Minerals
- Run a PHREEQC (Basic) Simulation
- Run a PHREEQC (Advanced) Simulation

6.1 Calculators and Converters

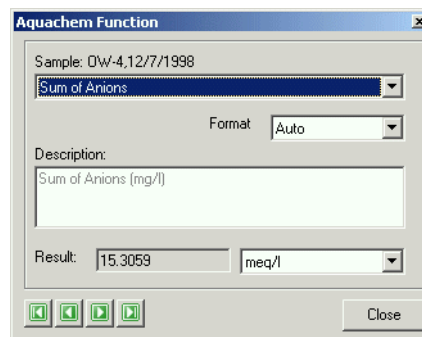
6.1.1 AquaChem Function

This tool allows you to quickly calculate any of the AquaChem built-in functions. Calculations can be performed without having to add/remove individual functions to the options of the Sample Details window.

To use this tool, first select a sample from your Active Samples list. Then select **Calculators > AquaChem Function** from the **Tools** menu, and a dialogue will appear as shown to the right side.

In this dialogue, you can see a **Description** of the function and the function units. By clicking on the down-arrow key, choose a function and then **Format**, and you will see the resulting value displayed in the **Result** field.

Use the  arrows to move through the samples.



The calculations can be customized in the **Calculations** section of the **Preferences** options dialogue under the **File** menu. In here, you may select which calculations should appear in the AquaChem functions (calculations) lists.

6.1.2 Decay Calculator

The **Decay Calculator** allows you to calculate the concentration of a contaminant at a particular point in time following a specified rate of decay, or the time to reach a specified concentration based, again, on that decay rate. AquaChem also allows you to calculate the decay rate of a chemical based on two or more concentration-time pairs. Degradation rate of a contaminant is usually given as a "half-life", which is the time required for the substance to be reduced to half of the initial mass. This tool can be applied only to organic chemical parameters in your database. The formula for the rate of decay is as follows:

$$C_0/C_1 = e^{-kt}$$

Where,

t = time

C₀ = initial concentration

C₁ = concentration after time t

When you access this tool, the following **Decay Calculator** dialogue will appear. The components of this dialogue are described below.

Sample

At the top of this dialogue under the **Sample** field, click on button and then double-click in **Pick a sample** list to select a sample for which you would like to run a degradation analysis. Selecting a sample from the sample list copies the concentration value for the current parameter to the initial concentration field **C(t=0)**.

Parameter

In the **Parameter** field, click on the button and select the organic parameter you would like to analyze (please ensure that your sample contains a value for this organic parameter). Note that the parameter list contains all database parameters, which have a corresponding record in the **Degradation Rates** table (available under **Tools > LookUp Tables**).

Half-Life

After selecting a parameter, the **Half-Life** field will be automatically filled in with the appropriate value. The combo box to the right of the **Half-Life** field provides options for the Half-Life environments (**Soil**, **Air**, **Groundwater**, and **Surface water**) for which degradation rates are specified in the database. If all rates are known for these environments the list will contain a **high**, **low**, and **average** half-life values for each environment. Effects such as dilution, transport or adsorption etc. are not taken into account with this tool.

AquaChem includes values for 335 organic chemicals. These are taken from the following Reference:

Howard, P.H. et al., 1991: *Handbook of Environmental Degradation Rates*, Lewis Publishers, 775 p.

This tool may also be used as a simple degradation calculator, without selecting samples or parameters. Simply enter a **Half-life** value manually, choose your **Problem Type**, enter an initial concentration, target concentration or time period and press **Calculate**.

AquaChem also allows you to estimate a chemical's half-life based on two or more time-concentration pairs.

Time Unit

Half-life values in the AquaChem database are saved as hours, however, you may convert these to days or years by selecting the appropriate **Time Unit** from the combo box.

Concentration Unit

Allows you to select the concentration units that will be used in the half-life calculations.

Problem Type

In the next line, you must specify the **Problem Type** for the degradation calculator. The **Decay Calculator** has three problem types available:

Type 1: Time to reach a specified concentration

Type 2: Concentration after a specified amount of time

Type 3: Estimate half life from given concentration values

NOTE: If problem type 1 is selected, and the current contaminant has an assigned guideline level in your AquaChem database, a combo box with available guideline levels will appear to the right of the target concentration field. You have the option to choose the desired guideline level, and the **C1** value will be automatically filled in for you.

A summary of the required data input is as follows:

Problem Type 1

In Problem Type 1 you can determine the time required for a contaminant to decay to a specified concentration.

C(t=0): initial concentration

C1: target concentration

Press **[Calculate]** and the **t(C1)** (time to reach target concentration) will be calculated.

The screenshot shows a software interface titled "t(C)". It contains three input fields: "C(t=0) = 500 ug/l", "C1 = 10 MCL ug/l", and "t(C1) = [blank] days".

Problem Type 2

In Problem Type 2 you can determine the concentration after a specified amount of time.

C(t=0): initial concentration

t1: target time

Press **[Calculate]** and the **C(t1)** (concentration after a specified time) will be calculated.

This calculator accounts only for degradation effect, using the following equation:

$$C(t) = C1 * e^{-kt}$$

where,

k = residence time

The screenshot shows a software interface titled "C(t)". It contains three input fields: "C(t=0) = 500 ug/l", "t1 = 10", and "C(t1) = 324 days".

Problem Type 3

In Problem Type 3 you can estimate half life from given concentration values.

Set the **Parameter** to "Custom"


Time: time of sampling

Conc.: concentration at sample time

Click **[Calculate]** to calculate the half-life as the slope of the regression line through the specified points. For every specified point, the estimated concentration by means of the linear regression line is displayed to verify the quality of the calculated values. At the left hand side of the screen, the estimated initial concentration, half-life, and the regression coefficient are displayed.

The screenshot shows a software interface titled "Estimate Halflife". It features a table with three columns: "Time", "Conc.", and "Est. Conc.". The table contains three rows of data: (1, 100, 100.1), (10, 75, 74.8), and (20, 54, 54.1). To the right of the table, there are three input fields: "C(t=0) = 104.2", "Half-life = 21.1", and "regr. coeff. -1.00".

Time	Conc.	Est. Conc.
1	100	100.1
10	75	74.8
20	54	54.1

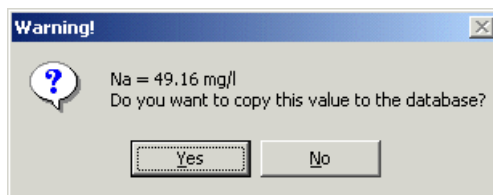
You can edit the concentration-time pairs at any time by double-clicking in the cell that contains the value you wish to edit. To remove the concentration-time pair highlight any cell in that pair and click the  button.

6.1.3 Find Missing Major Ion

If information for one of the major anions or cations (Na, Ca, Mg, Cl, HCO₃, SO₄) is missing from a sample, you can use this tool to estimate the concentration of the missing major ion. The concentration of the missing ion is calculated using the assumption that the total concentration of cations in meq/l is equal to the sum of anions. The entire difference of charges is assigned to the missing parameter.

NOTE: This tool is only available when you have a **Sample Details** window open and active.

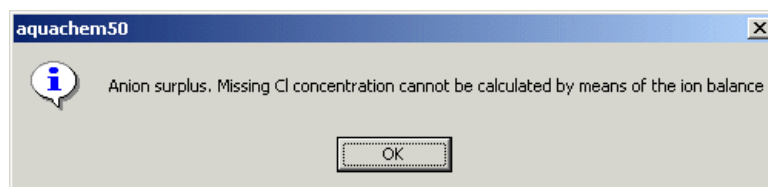
Open a sample, and select **Tools > Calculators > Find Missing Major Ion > Na** (for the example shown below).



After you have selected the desired missing parameter, AquaChem will calculate the missing value and display a confirmation message as shown on the right. Click **[Yes]** to accept the calculated value, or click **[No]** to reject the calculated value.

This tool does not work if more than one of the major cations or anions is missing in the analysis. In order to calculate the missing values for one of the major ions, each of the remaining major ions must be present.

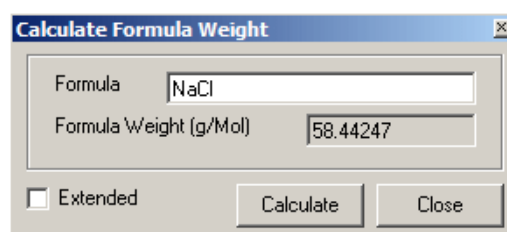
If there is an over-abundance of the missing ion's charge (e.g. Cl concentration is missing, the sample has already an anion excess) the following message will be displayed:



6.1.4 Formula Weight Calculator

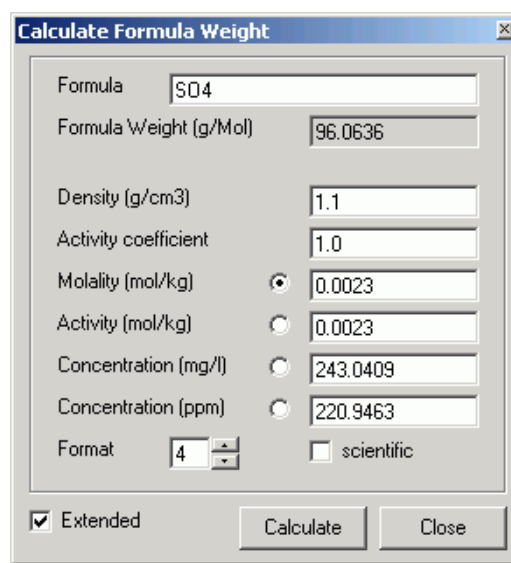
This tool allows you to quickly calculate formula weights for a user-specified chemical formula.

When you access this tool, the **Calculate Formula Weight** dialogue will appear. Simply enter the desired formula, and press **[Calculate]**.



Note that the **Formula** is case sensitive; therefore, ensure that you enter two-letter elements such as Silicon as Si and not SI. In addition, only one level of parentheses is taken into account. So, in case of a formula with two levels of parenthesis such as ((XY2)3)3AC, it will not be calculated correctly. In the standard display, simply the weight of the specified formula is calculated.

When checking the **Extended** checkbox, more options become available to convert different types of concentrations. This is convenient for converting PHREEQC results, expressed as mol/kg into mg/l. To do so, first enter the formula of the chemical. Then copy the PHREEQC molality value into the respective field, check the radio button and press the calculate button. The radio button tells the system that this field has been entered manually, and that all other concentration fields should be recalculated based on this value, the formula weight of the chemical, the density of the solution and the activity coefficient. this option may also be used



to convert mg/L concentrations into ppm, provided that the density of the solution is known.

6.1.5 Aggregate Samples

The aggregate samples tool allows you to create a new sample by aggregating existing samples. Parameters can be aggregated using various aggregation methods, including Minimum, Maximum, Median, Geometric Mean, Quantile and Custom.

When this option is selected from **Tools > Calculators > Aggregate Samples**, the Aggregate Samples dialog will appear on your screen.

Station	Date	Depth	Temp	pH	DO	TOC	TSS	Cond	TDS
MW-1	8/15/1992	5	14.8	7.15	9.2			1200	726
MW-1	6/1/1993	10	15.5	7.4	10.4			1170	1132
MW-1	6/15/1994	15	15	7.3				1470	1474
MW-1	7/30/1995	20	13	7.5				1295	1346
MW-1	7/28/1996	25	14.2	7.3				1270	1260
MW-1	6/15/1997	30	14	7				1225	1000
MW-1	8/1/1998	35	15	6.99				1310	1105
MW-2	8/1/1992	5	15.4	7.35				1140	1135
MW-2	6/5/1993	10	15.4	7.55				1190	1060
MW-2	6/12/1994	15	15.7	7.3				1300	1100
MW-2	7/21/1995	20	15.1	7.3				1230	1078
MW-2	7/24/1996	25	15	7.1				1240	1248
MW-2	6/1/1997	30	15.7	7.3				1100	1500
MW-2	7/24/1998	35	15.7	7.15				1060	1514
MW-3	8/8/1992	5	14	7.4				674	630
MW-3	6/8/1993	10	15	7.3				729	630
MW-3	6/15/1994	15	13	6.8				800	680
MW-3	7/25/1995	20	13	7.55				569	480
MW-3	8/2/1996	25	12.9	7.2				594	464
MW-3	6/6/1997	30	14.8	7.3				741	700

The Aggregate Samples dialog consists of two tabs: **Aggregation** and **Options**.

Under the **Aggregation** tab, each sample currently displayed in the Active Samples list is displayed in a table, along with the values for each analyzed parameter. The very last row in the table contains the aggregation value for each parameter.

The aggregation method can be selected from the **Aggregation** combobox, located directly above the table. When you select a new aggregation method, be sure to also select the **[Refresh]** button to repopulate the table with the new aggregation values.

The **Options** tab provides options for configuring a customized aggregation method. This tab is only applicable when the **Custom** option is selected under the **Aggregation** tab. The custom aggregation method allows you to mix two aggregation types. For example, average for major ions and geometric mean for trace methods. Choose the desired methods from the **Method 1** and **Method 2** combo boxes, and then associate each parameter with a aggregation method in the table below.

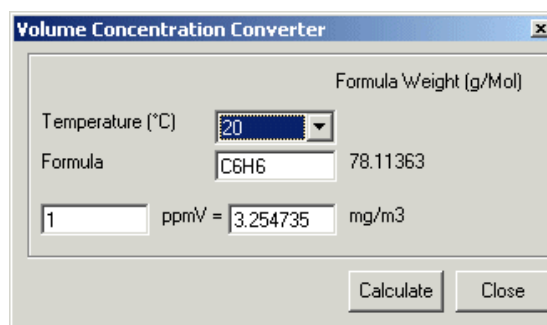
Once the desired aggregation method is selected, click the **[Save]** button, and chose whether to create a new sample, or to overwrite an existing sample with the aggregated values.

Please note that overwriting an existing sample with aggregated values cannot be undone.

6.1.6 Volume Concentration Converter

This converter allows you to quickly convert volume concentrations into mass concentrations for gas species, which are frequently reported in either of the two units.

When you access this tool, the **Volume Concentration Converter** dialogue will appear. Simply choose the observed ambient **Temperature** (0, 20, or 25°C), enter a **Formula** for an organic chemical, and enter the concentration. You may enter the concentration in either the **ppmV** or the **mg/m3** field.



Press **[Calculate]** and the missing concentration will be calculated.

In the vapor phase, one ppm by volume (ppmV) is on a volume per volume bases. For example, one ppmV of Benzene in the air means one part volume of benzene in one million parts volume of air space. To convert the ppmV into mass concentration units (mg/m3), the following formula can be used:

$$1 \text{ ppmV} = \text{FMW}/K$$

where,

FMW = Formula Weight (g/mol)

K = is a temperature dependant coefficient (molar gas volume).

K = 22.4 l/mol at 0°C,

K = 24.05 l/mol at 20°C, and

K = 24.5 l/mol at 25°C.

Example:

To convert 1ppmV Benzene to mg/m3 at 20°C:

$$\begin{aligned} \text{Mass Benzene mg/m3} &= 1\text{ppmV Benzene} * [(78 \text{ g/mol}) / (24.05 \text{ l/mol})] \\ &= 3.24 \text{ g/L} \end{aligned}$$



$$= 3.24 \text{ mg/m}^3$$

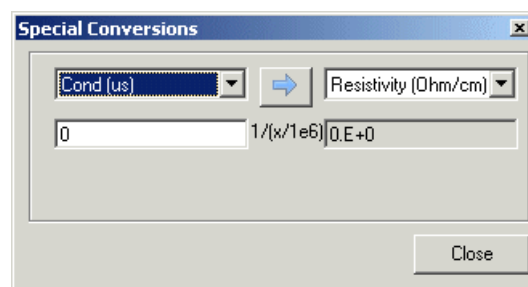
NOTE: The formula is case sensitive, therefore, ensure that you enter two-letter elements such as Silicon as Si and not SI. In addition, only one level of parentheses is taken into account (i.e. a formula such as ((XY2)3)3AC will not be calculated correctly).

6.1.7 Special Conversions

This tool provides conversions for numerous non-linear geochemical calculations. You may convert values from one measurement unit to another for measurement units such as pe - Eh, Alkalinity - HCO₃, and Conductivity (us) - Resistivity (Ohm/cm).

When you access this tool, the **Special Conversions** dialogue will appear.

Click on the  button and choose a category from the list, enter the start value, and any other required variables (temperature, density), and press  to convert the value.




The available conversion categories include:

Conductivity  Resistivity

where,

$$\text{Cond} = 1/(\text{Resistivity}/1\text{E}+6)$$

Electrical conductivity at a different temperature: Cond(T1)  Cond(T2).

It requires you to enter a conversion temperature. The conductivity is always normalized to a specific temperature which is dependant on the probe. If you want to enter conductivity data to your database, and the temperature for the probe is different (e.g. 25°C) from the one in the database (20°C) then the data has to be converted before being adding to the database.

This calculation is done in two steps:

- First, calculate the conductivity at standard temperature,
- Second, calculate the conductivity at the specified temperature (Temp2).

Examples:

Conductivity

$$\text{Cond}(25) = \text{Cond}(\text{Temp1}) / (1 + 0.0198 * (\text{Temp1} - 25))$$

$$\text{Cond}(\text{Temp2}) = \text{Cond}(25) * (1 + 0.0198 * (\text{Temp2} - 25))$$

Redox potential

$$\text{pe} \rightarrow \text{Eh}$$

It requires you to enter an observation temperature.

$$\text{pe}(\text{Eh}): \text{pe} = \text{Eh} * F / 2.303 / R / T$$

$$\text{Eh}(\text{pe}): \text{Eh} = \text{pe} * 2.303 * RT / F$$

where,

$$R = 8.314 \quad (\text{Gas constant})$$

$$F = 96485 \quad (\text{Faraday constant})$$

Reference: C.A.J. Appelo, 1996: Geochemistry, Groundwater and Pollution, Balkema Rotterdam, p. 246.

Alkalinity calculations

Convert between:

mg/L HCO₃,

meq/L,

Alkalinity (f*), where * means French degrees,

Alkalinity (g*), where * means German degrees

mg/L CaCO₃.

Reference: John D. Hem. Study and Interpretation of the Chemical Characteristics of Natural Water, USGS Water supply paper 2254, p. 158.

Other conversions

ppm \rightarrow mg/L: Requires you to enter a density.

$$\text{mg/L} = \text{ppm} / \text{Density}$$

mg/L \rightarrow ppm: Requires you to enter a density

$$\text{ppm} = \text{mg/L} * \text{Density}$$

6.1.8 Species Converter

This tool allows you to convert any aqueous species into a different form. When you access this tool, the **Species Converter** dialogue will appear as shown below:

		Formula Weight (g/Mol)
Master Element	Si	28.0855
Source Species	SiO ₂	60.084
Target Species	H ₂ SiO ₃	78.100
Conversion Factor	1.300	

10 mg/l SiO₂ = 12.998 mg/l H₂SiO₃

- Select a **Master Element** (Parameter) in which you are interested (common examples include nitrogen, phosphorus, silica, etc.).
- Type in a **Source Species**.
- Type in a **Target Species**.
- Enter a concentration of the source species in the lower left corner of the dialogue.
- The **Conversion Factor** is calculated based on the formula weights of the two species: the **Target species** divided by the **Source Species**.
- Press **[Calculate]** and the concentration for the target species will be calculated.

This tool is practical for expressing a measured amount of a parameter as different aqueous species when expressed in mg/L. For example, you may receive your Silica results as mg/L Si, and your database requires mg/L SiO₂ or H₂SiO₃. For this you need to calculate the amount of SiO₂ that holds the same amount of Silica as given in the lab analysis. Nitrogen species such as nitrate and nitrite may are commonly expressed as N or as their actual formula (NO₃ or NO₃) and constitute another good example where the above calculation may be useful.

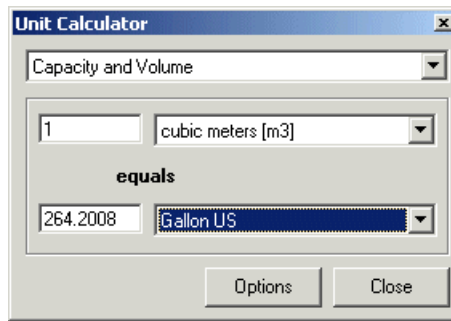
The data entered in the **Species Converter** dialogue (shown above) depicts an example for converting Si species.


NOTE: The Formula is case sensitive; therefore, ensure that you enter two-letter elements such as Silicon as Si and not SI. In addition, only one level of brackets is taken into account (i.e. a formula such as ((XY₂)₃)₃AC will not be calculated correctly).

6.1.9 Unit Calculator

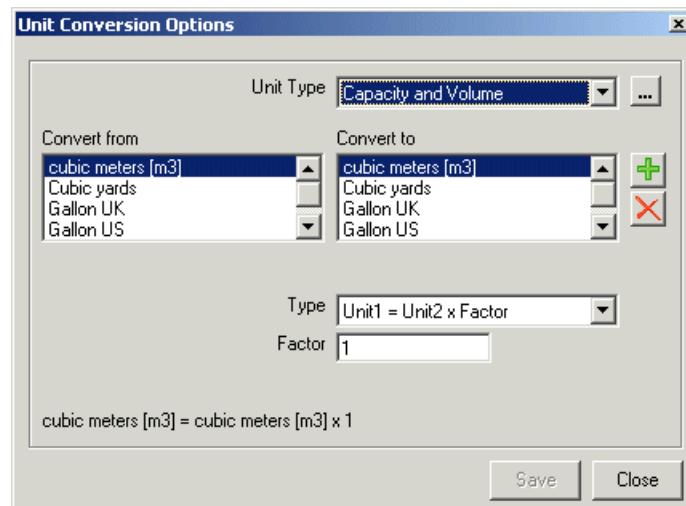
This tool performs basic unit conversions for length, time, volume, density, mass, etc.



When you access this tool, the **Unit Calculator** dialogue will appear as shown below.



Click on the  button and select the desired category from the combo box at the top of the dialogue, enter the value and units in the fields below, and the new value will be automatically calculated.

In addition, you can modify the existing conversion factors, or create new ones. Press the **[Options]** button, and the following dialogue will be loaded.



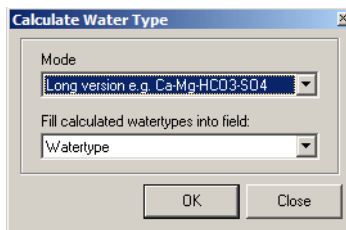
Select the **Unit Type** from the combo list (at the top), or press the  button to load a list of available unit types, and options to create a new unit type. The list below will now show all conversions available for this unit type, and the conversion factor associated with each. You may enter or edit all factors, or create new conversions using the  button. Then enter the appropriate conversion type and factor for this new conversion.

Once you are finished, press **[Save]** to save any changes you made to the conversions.

Press **[Close]** to return to the **Unit Calculator** tool.

6.1.10 Calculate facies

This command calculates the **Watertype** (water facies) expression for all highlighted samples. The following dialogue allows you to chose the algorithm to be used in the calculation and the field where the calculated value will be placed:



The **Mode** lets the user select whether he wants to use the long facies name, which is based on the ions that contribute more than 10% (e.g. Ca-Mg-Cl-SO₄), or the short mode, listing just the most frequent anion and cation.

The **Fill calculated waternypes into field** combo box allows you to specify the target field where the value should be stored. Every sample must contain a field which contains the watertype and which is automatically recalculated if any changes to chemical concentrations occur, however the user may specify a second field, holding the water type in short notation. This can be useful when creating symbols based on waternypes. To implement this, use he **Calculate facies** command and select the **Short version** in the **Mode** combo box. Then, specify a field other than the official water facies field (where you want to keep the more precise long notation). Next, go to **Symbols and Lines** dialogue and create a new symbol group called Water type. Use the automatic function based on the short watertype field to create a new symbol for every distinct value of water facies (Ca-HCO₃, Ca-SO₄, Na-Cl et.c)

In most cases the procedure of calculating the long water facies is not necessary since the facies expression is calculated automatically when importing data or when entering or modifying data in the **Sample Details** window.

However, in some cases the chemical composition of a sample might change without having this change reflected by the water facies. For example, the facies expression is not updated for all samples entered or modified in the **Table View**. Additionally there are a few commands allowing to change the chemistry of a sample without passing by the sample screen, such as using the **Replace/Multiply** function, marking extreme values as outliers using the **Outlier Tests**, or calculating the HCO₃ and CO₃ concentrations from the measured alkalinity. For these cases, the water facies may be synchronized manually with the chemical composition using the **Calculate facies** command.

As outlined in section 3.11, recalculation of the water facies expression may be necessary, after modifying the way for calculating the water facies expression.

6.1.11 Corrosion & Scaling

Corrosion and scaling of well equipment count among the most common concerns in water management. Various methods have been developed to predict the tendency of the ground or surface water to precipitate minerals or to corrode metal pipes. The most commonly used method is the Langelier Saturation Index LSI. The Ryznar Stability Index RSI is closely related to the LSI. Both indexes are based on the following parameters: Temperature, pH, TDS, alkalinity and calcium. Various ranges and associated risks of scaling or corrosion have been suggested and AquaChem follows the ranges outlined below in table 1 and 2. Ranges and comments associated with the ranges may be changed in the **File>Database>Ranges** section.

Table 1: Ryznar stability index

Value	Interpretation
< 5	Heavy scaling
5 - 6	Mild scaling
6 - 7	Little scale or corrosion
7 - 7.5	Corrosion significant
7.5 - 9	Heavy Corrosion
>9	Corrosion intolerable

Table 2: Langelier Saturation Index

Value	Interpretation
<-2	Serious corrosion, CaCO ₃ dissolution
-0.5 - -2	Slightly corrosive but non-scale forming
-0.5 - 0.5	Balanced but pitting corrosion possible
0.5 - 2	Slightly scale forming and corrosive
>2	Scale forming but non corrosive

LSI and RSI are empirical methods based on a limited number of parameters and taking in account the precipitation of calcite only. They are useful in that they can be easily calculated and interpreted by means of tables such as table 1 and 2 and cover the most common scaling thread (calcite). However we suggest using the scaling and corrosion calculator only for a preliminary overview and base corrosion and scaling predictions on a PHREEQC simulation. PHREEQC will provide more reliable results in that it is based on a thermodynamic dataset and includes more corrosion/scaling relevant minerals and parameters such as Eh, Mn, Fe, iron hydroxides, etc.

If you want to calculate the corrosion and scaling indexes of a sample in the database, select the sample first, then select **Tools>Calculators>Corrosion and Scaling** menu command. The calculator will display as shown, and will be initialized with the values for this sample. If you launch the calculator without selecting a sample first, then the screen will open with empty fields which you must fill. When the fields are ready, press the **[Calculate]** button. You may test how the LSI and RSI react to changes of one or several parameter (e.g. for rising temperature) by changing the field values.

Note that for if the calculator has been initialized from a existing database sample and the Calcite saturation index has been calculated. This value will be displayed at the bottom of the screen, which may help to interpret the RSI and LSI. If there is a major discrepancy between the empirical indices and the modeled saturation index, you should proceed with a PHREEQC simulation on this sample. The PHREEQC Calcite saturation index value reflects the sample stored in the database. The comparison of calcite scaling indices such as Langelier and Ryznar to the PHREEQC saturation indices should only be verified as long as the original input parameters (temperature, alkalinity, calcium etc.) have not been changed. Once these values are modified, the corrosion and scaling indices will be recalculated based on the new input whereas the PHREEQC saturation index will still be based on the initial conditions.

Parameter	
pH	7.3
Temp (°C)	15.7
Alkalinity (mg/l CaCO ₃)	119.7462
Ca (mg/l)	124
TDS (mg/l)	0
SO ₄ (mg/l)	299
Cl (mg/l)	361

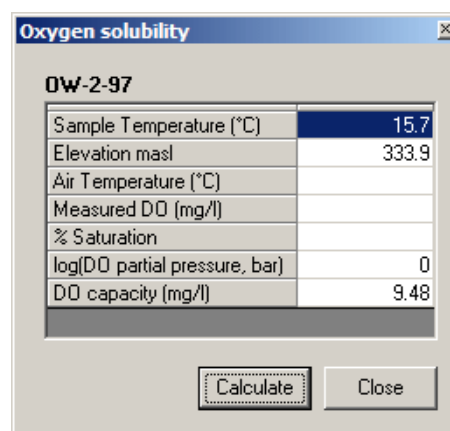
Index	
Saturation pH	7.30
Langelier Saturation Index (LSI)	0.00
Ryznar Stability Index (RSI)	7.29
Puckorius Scaling Index (PSI)	7.01
Larson-Skold Index	5.55

Calculate Close

6.1.12 Oxygen Solubility

This function allows you to calculate the oxygen solubility of a given sample as a function of elevation and temperature.

It is accessed from the main menu via **Tools > Calculators > Oxygen Solubility**



DW-2-97	
Sample Temperature (°C)	15.7
Elevation masl	333.9
Air Temperature (°C)	
Measured DO (mg/l)	
% Saturation	
log(DO partial pressure, bar)	0
DO capacity (mg/l)	9.48

Calculate Close

Dissolved oxygen saturation values are calculated using the American Society of Civil Engineers' formula (American Society of Civil Engineering Committee on Sanitary Engineering Research, 1960):

$$DO_{\text{sat}} = 14.652 - 0.41022T + 0.00799107T^2 - 0.000077774T^3$$

where DO_{sat} = dissolved oxygen saturation concentration, mg/L

T = water temperature, degrees Celsius

This formula represents saturation values for distilled water at sea level pressure. The Dissolved oxygen saturation concentrations generated by the formula must be corrected for elevation above mean sea level due to differences in air pressure, as well as differences caused by air temperature changes. The correction factor is calculated by the Oxygen Solubility Calculator using the following:

$$f = \frac{2116.8 - (0.08 - 0.000115A)E}{2116.8}$$

Reference: Lin, S., 2001: *Water and wastewater calculations manual*. McGraw-Hill, New York. pp. 6-8.

6.1.13 UTM Conversion

This function allows you to calculate geographic coordinates from projected (UTM) coordinates, and vice versa, for stations in your database. Geographic and UTM station coordinates are stored in separate columns in the aquachem database and can be included in exported station data, e.g., .KML, .SHP etc.

To use this tool, select **Tools > Calculators > UTM Calculator** from the main menu.

The UTM Conversion dialog will appear on your screen.

UTM conversion

Mode: Geographic to UTM

Datum: NAD83/WGS84

StationID	X	Y	Coord_Long	Coord_Lat
MW-1	535250.2	4814315	0	0
MW-2	535535.5	4814905	0	0
MW-3	536668.1	4814036	0	0
MW-4	536720.7	4814826	0	0

Calculate Close

Select the desired conversion type from the **Mode** combo box. If you wish to calculate UTM coordinates from Geographic coordinates, select the Geographic to UTM option. If you wish to calculate Geographic coordinates from UTM coordinates, select the UTM to Lat/Long option.

Geographic to UTM

When this option is selected, you must also specify the associated datum of the existing geographic coordinates.

UTM to Lat/Long

When this option is selected you must also specify the associated UTM direction and zone of the coordinates.

Location

Direction: N W

Zone: 4

Click the **Calculate** button to calculate the coordinates. Please note that any existing coordinates will be overwritten with the new values.

6.1.14 Retardation

Retardation factor (Rf) for a particular chemical species is the ratio of solution velocity and species velocity or the ratio between the rate of groundwater movement and the rate of contaminant movement. Rf is a required input for transport modules.

$$\text{Retardation Factor} = 1 + (\rho_b/\theta) (K_d)$$

Where:
 ρ_b = dry bulk mass density of the soil, gm/cc
 θ = Porosity
 K_d = distribution coefficient for the solute with the soil, ml/g
C = Concentration of solute
C* = Concentration of solute on solid

Retardation Calculator

Calculation | Options

Parameter	Value
Grain density (g/cm^3)	2.7
Bulk density (g/cm^3)	2.16
Porosity	0.2
Organic fraction	0.01
Log Kow (octanol-water partitioning coefficient)	2.13
Koc from	log(koc)=a*log(kow) + b
Log Koc (organic carbon-water partitioning coefficient)	1.76
Kd (distribution coefficient, mL/g)	0.58
R (retardation coefficient)	7.21

CalculateClose

6.2 QA/QC

6.2.1 Reliability Check

This **Reliability Check** report helps you confirm the validity of the measured sample data. When you select this report, the following **Reliability Check** window will appear:

Sample Summary Report			
Sample ID	MIV-1-03		
Sample Date	6/1/1993		
Station	MIV-1		
Location	silty-gravel		
Geology	Ca-Na-SO4-Cl-HCO3		
Water type	15.5		
Temperature (°C)	1170 uS/cm		
pH			
Conductivity			
Sum of Anions	12.90578	meq/L	
Sum of Cations	55.05045	meq/L	
Balance	62.01738	%	
Total dissolved solids	1703.231	mg/L	
Total hardness	0	mg/L CaCO3	
Alkalinity	0	mg/L CaCO3	
Major ion composition	mg/L	mmol/L	meq/L
Na	60	2.957837	2.957837
K	1.4	3.580718E-02	3.580718E-02
Ca	1000	24.95135	49.90269
Mg	26	1.069739	2.139477
Cl	121	3.412989	3.412989
SO4	225	3.355417	6.710833
NO3	<0.1	+1.612903E-	+1.612903E-
HCO3	158.6	2.59927	2.59927
Ratios	Sample	mg/L	Standard Seawater
Ca/Mg	38.46154	23.32471	0.319
Ca/SO4	3.076923	7.370244	0.152
Na/Cl	0.5619825	0.8666391	0.656
Cl/Br			287

This report provides a number of checks which can lend insight as to the reliability of the water sample analysis. If the analysis value passes the test, then a **Pass** will be displayed in the **Result** column; if not, then a **Fail** will be displayed. Each analysis is explained in the table below:

Available Tests in the Reliability Check Report

Test	Attention Value	Comments
Balance (C-A)/(C+A)*100	<5%	The solution must be electrically neutral or within ± 5 : sum of cations = sum of anions.
TDS: [(Entered-calculated)/Entered]*100	<5%	Calculated TDS = sum ions (mg/L) + SiO2 must be similar to measured dry residue.
TDS: [(Entered -TDS180° calculated)/Entered]*100	<5%	Calculated TDS = sum ions (mg/L) + SiO2 - 0.5082 * bicarbonate. Must be similar to measured dry residue at 180°.
TDS Entered/Conductivity	0.55 < x < 0.75	There is a linear relationship between TDS and conductivity within a range of 0.55 to 0.75.
Conductivity/Sum MEQ Cations	90 < x < 110	There is a linear relationship between conductivity and Sum of Cations within a range of 90 - 110.
K ⁺ /[Na ⁺ + K ⁺] meq/l	<20%	Na >>K

Available Tests in the Reliability Check Report

$Mg^{++}/[Ca^{++} + Mg^{++}]$ meq/l	<40%	Mg is mainly provided by the dissolution of dolomite resulting in a Ca/Mg ratio of 1:1. However calcium has various additional sources such as the dissolution of gypsum or carbonate.
$Ca^{++}/[Ca^{++}+SO_4^{--}]$	>50%	Most SO_4 concentration can be attributed to the dissolution of gypsum. Therefore the Ca/ SO_4 ratio must be 1:1 or lower, if some Ca is also provided by the dissolution of carbonate.
$Na^+/(Na^+ + Cl^-)$	>50%	Chloride is mainly provided by the dissolution of Halite (NaCl). Therefore the ratio Na/Cl is 1 or higher, if some Sodium is added to the solution by the solution of silicates or by ion exchange.
SI Calcite	<0.2	In a normal groundwater, the saturation index of rock forming and reactive minerals may be assumed to be close to 0 (saturation) or below. Any supersaturation, indicated by $SI > 0$, will be rapidly compensated by the precipitation of the respective mineral. A significant calculated supersaturation of calcite normally indicates an error in pH measurement or postsampling addition of calcium due to dissolution of carbonate particles in the sampling vessel.
Test for results > solubility		The solubility of a chemical parameter may be defined in the database dialogue. If a measured result exceeds the theoretical solubility, the respective concentration should be checked
Test for results outside natural occurrence range		The natural occurrence range of a chemical parameter may be defined in the database screen using the range dialogue. If a measured result exceeds the defined range, the respective concentration should be checked

NOTE: Some attention values are displayed as “acceptable ranges”. This means that the Analysis value must be within this range. The analysis (calculated) values can be positive or negative. If the analysis values are outside this range (either positively or negatively) then the value will not “pass” this check, and the report will display a “Fail”, indicating that the sample did not pass this analysis check.

If the calculated values are not within the attention values (i.e. the Result is **Fail**), then this does not necessarily signify an error; it does mean however that there should be an explanation for the value. For example, if $Na/Cl < 1$ then the explanation could be that some Na has been removed from the solution by $Na > Ca$ exchange: this should be

confirmed by a suitable aquifer geology (exchange friendly minerals such as Ca-rich clay). A ratio of $\text{Ca}/\text{SO}_4 < 1$ can be explained by the dissolution of ferrous minerals such as pyrite. This effect is often observed in mine tailings. However, normal groundwater samples should fulfill all the criteria mentioned above.

The scroll arrows in the lower right corner of the **Reliability Check** window can be used to produce a report for other samples in your database.

For more details please see the reference below:

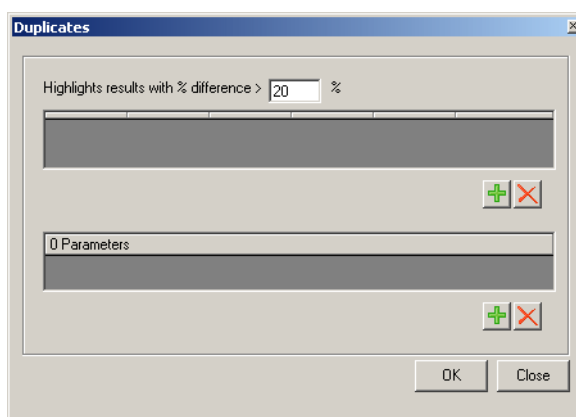
Hounslow, A.W. 1995. *Water Quality Data – Analysis and Interpretation*. CRC Press LLC. p.75.

6.2.2 Compare Duplicates

Field duplicates are a common quality control method for water quality data. A field duplicate is a duplicate sample, collected by the same team or by another sampler or team at the same place at the same time. It is used to estimate sampling and laboratory analysis precision.


In order to compare duplicates, duplicate samples must have an assigned duplicate code using the **Find Duplicates** utility (“Find Duplicates” on page 348).

To compare duplicates, select **Tools > QAQC > Compare Duplicates**, from the main menu. The following dialog will appear on your screen.




The Duplicates dialog allows you to specify which samples and parameters to include in the comparison.

Selecting Duplicate Samples

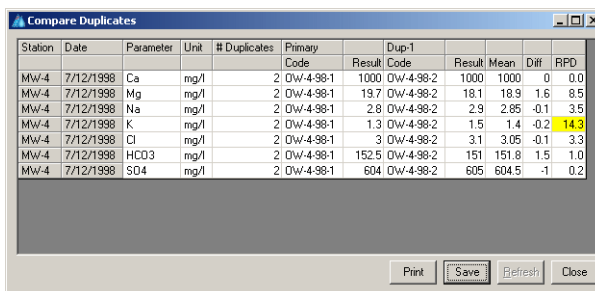
To specify which samples to include in the comparison, select the top  button. A list will appear displaying all samples in the database that have been assigned a duplicate code. Select the desired samples for comparison and then click the **[Assign]** button.

Selecting Parameters

To specify which parameters to include in the comparison, select the bottom  button. A list of parameters will appear displaying all measured parameters. Select the desired parameters for comparison and then click the **[Select]** button.

In the Duplicates dialog, select the **[OK]** button to generate the report.

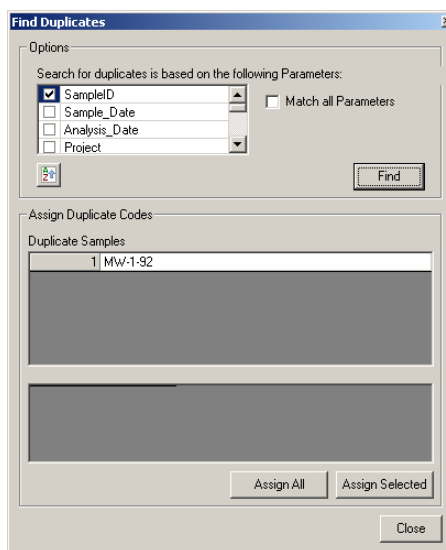
A report will appear on your screen with the a sample comparison, including the relative percent difference value.



Station	Date	Parameter	Unit	# Duplicates	Primary Code	Dup-1 Code	Result	Mean	Diff	RPD
MW-4	7/12/1998	Ca	mg/l	2	DW-4-98-1	1000 DW-4-98-2	1000	1000	0	0.0
MW-4	7/12/1998	Mg	mg/l	2	DW-4-98-1	19.7 DW-4-98-2	18.1	18.9	1.6	8.5
MW-4	7/12/1998	Na	mg/l	2	DW-4-98-1	2.8 DW-4-98-2	2.9	2.85	-0.1	3.5
MW-4	7/12/1998	K	mg/l	2	DW-4-98-1	1.3 DW-4-98-2	1.5	1.4	-0.2	14.3
MW-4	7/12/1998	Cl	mg/l	2	DW-4-98-1	3 DW-4-98-2	3.1	3.05	-0.1	3.3
MW-4	7/12/1998	HCO3	mg/l	2	DW-4-98-1	152.5 DW-4-98-2	151	151.8	1.5	1.0
MW-4	7/12/1998	SO4	mg/l	2	DW-4-98-1	604 DW-4-98-2	605	604.5	-1	0.2

6.2.3 Find Duplicates

The **Find Duplicates** utility allows you to find duplicate samples and assign duplicate codes:



Finding Duplicate Samples

Choose one or more parameters from the list of available parameters by placing a check mark in the appropriate box(es). Only Sample Description and Station Description parameters are available. To select all parameters, place a checkmark beside **Match all parameters**. Then click the **[Find]** button.

AquaChem will then run a query on the samples in the **Active Samples** list and locate all samples for which the values of the set of specified parameters are identical. Common parameters for this utility are sampleID and date.

The results will then be shown in the lower half of this dialogue.

Assigning Duplicate Codes

In order to compare duplicate samples using relative percent difference (RPD) analysis and reporting (see “Compare Duplicates” on page 347), each duplicate sample must be assigned a duplicate code. The duplicate code is stored in the Duplicate_ID field for each sample.

Note: Samples that have not been assigned a duplicate code will not be available for sample comparison using RPD analysis.

To assign a duplicate code to samples,

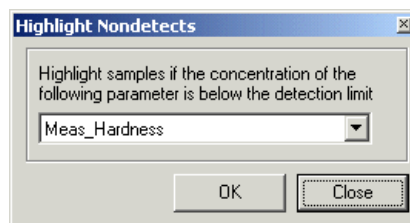
- Select the desired samples from the **Duplicate Samples** list (located at the bottom of the **Find Duplicates** dialog.)

- Select the **Assign Selected** button to assign the duplicate code to each selected samples.

To assign a duplicate code to all samples in the duplicate samples list, select the **[Assign All]** button

6.2.4 Highlight Nondetects

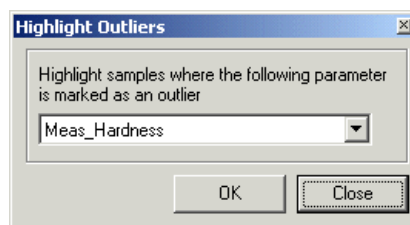
Selecting this option will load the following dialogue:



Selecting the appropriate parameter and clicking **[OK]** will highlight all samples that have non-detect value for that parameter.

6.2.5 Highlight Outliers

Selecting this option will load the following dialogue:



Selecting the appropriate parameter and clicking **[OK]** will highlight all samples that have that parameter marked as an outlier.

6.2.6 Highlight Duplicates

Selecting this option will highlight all samples that have a **Duplicate_ID** value entered. The Duplicate_ID is entered in the top portion of the Sample Details window.

6.3 LookUp Tables

The **Look-Up Tables** menu displays the available AquaChem tables containing useful hydrochemical information. Look-Up tables allow you to store and retrieve any useful information from within the AquaChem environment. The tables are created using MS Access, and are saved in the Aquachem.mdb database. These new tables can then be displayed and searched in Aquachem.

NOTE: You need the MS Access environment in order to create or import tables you want to make accessible in this way. AquaChem only allows you to visualize, search and edit the data.

The first three tables are used by AquaChem for internal calculations, and can not be modified or removed.

- Degradation Rates
- PHREEQC Phases
- Periodic Table
- Water Standards
- Time Series

The remaining three tables are examples of user-defined tables:

- Links
- Organic Compounds
- Preservation Methods

These tables can be removed by selecting **File** from the main menu and then **Database\Lookup Tables** dialogue. These tables are used for lookup purposes and are read-only. You may create up to ten user-defined tables. All other tables may be defined, provided you have MS Access2000 development environment.

The **Demo** database includes two example Look Up Tables, which are meant for demonstration purposes only and do not represent a complete dataset. They provide an example of useful data which can be used in an AquaChem project. You may create your own tables which may be used for a specific study, and bring these tables into AquaChem.

For more details on creating user-specified LookUp tables in AquaChem, please refer to the Lookup Tables in Chapter 3.

6.3.1 Degradation Rates

This table provides a list of **Degradation Rates** for organic compounds. Half-life values are available for various phases (Soil, Air, Surface water, and Groundwater). All half-life values are displayed in hours. The data from this table is used for the **Decay Calculator**. These degradation rates are from the following reference:

Howard, P.H. et. al. 1991. *Handbook of Environmental Degradation Rates*, Lewis Publishers, 775 p.

6.3.2 PHREEQC Phases

The minerals list is read from the PHREEQC thermodynamic database file. If no valid thermodynamic database is specified, the **List of Minerals** will be empty. This table displays PHREEQC phase names, formula, and molecular weights.

6.3.3 Periodic Table

This table provides information for all elements in the **Periodic Table**. This includes: Name, Formula, Mass, Valence, CAS #, and Group Name.

6.3.4 Water Standards

This table contains all of the water quality standards and guidelines defined in the project database. Use the comboboxes located at the top of the window to show specific guidelines, levels and parameters.

Water Standards

Guideline

WHO

Level

MCL



Standard

CAI

Type	Guideline	Level	Param (orig)	Unit (orig)	CASNo	Lower (orig)	Upper (orig)	Year	Lower(aqc)	Upper(aqc)	Comment
C	WHO	MCL	1,1-Dichloroethene	mg/L	75-35-4		0.03	0			
C	WHO	MCL	1,2-Dibromo-3-chloropropane	mg/L	96-12-8		0.001	0			
C	WHO	MCL	1,2-Dibromoethane	mg/L	106-93-4		0.0004	0			
C	WHO	MCL	1,2-Dichlorobenzene	mg/L	95-50-1		1	0			
C	WHO	MCL	1,2-Dichloroethane	mg/L	107-06-2		0.03	0			
C	WHO	MCL	1,2-Dichloroethene	mg/L	540-59-0		0.05	0			
C	WHO	MCL	1,2-Dichloropropane	mg/L	78-87-5		0.04	0			
C	WHO	MCL	1,3-Dichloropropene	mg/L	542-75-6		0.02	0			
C	WHO	MCL	1,4-Dichlorobenzene	mg/L	106-46-7		0.3	0			
C	WHO	MCL	2,4,5-Trichlorophenoxy acetic acid	mg/L	93-76-5		0.009	0			
C	WHO	MCL	2,4,6-Trichlorophenol	mg/L	88-06-2		0.2	0			
C	WHO	MCL	2,4-D (2,4-dichlorophenoxyacetic acid)	mg/L	94-75-7		0.03	0			
C	WHO	MCL	2,4-DB	mg/L	94-82-6		0.09	0			
C	WHO	MCL	Acrylamide	mg/L	79-06-1		0.0005	0			
C	WHO	MCL	Alachlor	mg/L	15972-60-8		0.02	0			
C	WHO	MCL	Aldicarb	mg/L	116-06-3		0.01	0			
C	WHO	MCL	Aldrin	mg/L	309-00-2		0.00003	0			
C	WHO	MCL	Antimony	mg/L	7440-36-0		0.02	0			
C	WHO	MCL	Arsenic	mg/L	7440-38-2		0.01	0		10	
C	WHO	MCL	Atrazine	mg/L	1912-24-9		0.002	0			
C	WHO	MCL	Barium	mg/L	7440-39-3		0.7	0		0.7	
C	WHO	MCL	Benzene	mg/L	71-43-2		0.01	0		10	
C	WHO	MCL	Benzo[a]pyrene	mg/L	50-32-8		0.0007	0			
C	WHO	MCL	Boron	mg/L	7440-42-8		0.5	0		0.5	
C	WHO	MCL	Bromate	mg/L	15541-45-4		0.01	0			
C	WHO	MCL	Bromodichloromethane	mg/L	75-27-4		0.06	0			
C	WHO	MCL	Bromofom	mg/L	75-25-2		0.1	0			
C	WHO	MCL	Cadmium	mg/L	7440-43-9		0.003	0		3	
C	WHO	MCL	Carbofuran	mg/L	1563-66-2		0.007	0			
C	WHO	MCL	Carbon tetrachloride	mg/L	56-23-5		0.004	0			
C	WHO	MCL	Chloralhydrate	mg/L	302-17-0		0.01	0			
C	WHO	MCL	Chlorate	mg/L	7790-93-4		0.7	0			
C	WHO	MCL	Chlordane	mg/L	57-74-9		0.0002	0			
C	WHO	MCL	Chlorine	mg/L	7782-50-5		5	0			
C	WHO	MCL	Chlorite	mg/L	7758-19-2		0.7	0			
C	WHO	MCL	Chloroform	mg/L	67-66-3		0.2	0			
C	WHO	MCL	Chlorotoluron	mg/L	15545-48-9		0.03	0			
C	WHO	MCL	Chlorpyrifos	mg/L	2921-88-2		0.03	0			
C	WHO	MCL	Chromium	mg/L	7440-47-3		0.05	0		50	

Close

6.3.5 Time Series

This table contains all imported time series data. Use the combo box at the top to select the desired time series to show in the table below. You may add and delete records from the table using the  **Add** and  **Delete** button, located in the bottom-left corner of the dialog.

6.3.6 Links

This table contains links to several commonly used websites such as IUPAC Homepage and the PHREEQC Homepage. When double clicking the URL field, the respective web address will be opened in your browser or in case where this field contains a file name, the file is opened with its respective application (E.g. MS Excel or MS Word). This table is therefore well suited to created your list of favorites within AquaChem in order to access quickly often used resources on the internet, your local network or you local machine.

6.3.7 Organic Compounds

This table includes details on organic compounds. This includes: Name, CAS#, Formula, Specific Gravity, Solubility, K, Vapor Pressure, and HR.

6.3.8 Preservation Methods

This table lists some common sample preservation techniques.

6.4 Modeling

The **Modeling** option under the **Tools** menu provides links to the PHREEQC geochemical modeling utilities.

PHREEQC provides six options for geochemical calculations:

Option 1: Calculation of the aqueous equilibrium (activities and saturation indices) based on a sample analysis

Option 2: Calculate pH based on a sample analysis

Option 3: Calculate Eh based on a sample analysis

Option 4: Calculate bicarbonate and carbonate concentration based on the measured total alkalinity and pH

Option 5: Equilibrate with minerals present in the sample

Option 6: Basic forward modeling, such as mixing samples, adding minerals or chemicals to a solution, or raising temperature

Option 7: Advanced modeling such as transport calculations, inverse modeling, etc.

Option 1: Calculate Saturation Indices and Activities

AquaChem allows you to quickly calculate saturation indices and activities for the modeled parameters listed in your database; the results are automatically saved back to your database, provided that the fields exist in the database. For example, to read back the calcite saturation index calculation, you need a field SI_Calcite specified in the **Database Options, Modeled Parameters** section.

Option 2: Calculate pH

AquaChem allows you to use PHREEQC to calculate a pH value for your sample. This can be used in an instance where you do not have a pH value for your sample, or you would like to compare to a theoretical pH if the sample was exactly in equilibrium with a specified carbonate mineral (e.g. calcite or dolomite).

Option 3: Calculate Eh

AquaChem allows you to use PHREEQC to calculate the Eh value for your sample. The calculation is based on the available redox couples for the selected sample. The sample must include concentrations for at least 1 redox couple, e.g. Fe²⁺/Fe³⁺, Mn²⁺/Mn³⁺, etc.

Option 4: Alk > HCO₃, CO₃

Aquachem allows you to calculate the concentrations of the carbonate species HCO₃⁻, CO₃²⁻ as well as OH⁻ (if defined as a parameter in AquaChem) based on the alkalinity measurement. Note that this option is only active, if “Measured Alkalinity” has been matched to a database parameter in the **File > Database Aliases** dialogue.

Option 5: Equilibrate with Minerals

This option allows you to simulate how much mineral would dissolve or precipitate, if a given solution was brought into equilibrium with one or several minerals. You may change the temperature to simulate heating or cooling of the solution and you may also simulate evaporation. The following examples illustrate two classic applications:

- Calcite saturated water is pumped and its temperature raises from 12 to 25 degrees in the piping system. How much precipitated calcite can be expected if equilibrium is maintained?
- An initial solution evaporates in a pond. How much gypsum precipitates if the residual water is one tenth of the original volume?

Option 6: PHREEQC (Basic)

AquaChem also allows you to do basic modeling with the PHREEQC interface included with AquaChem. To do so, you must use the **PHREEQC (Basic)** option. This option allows you to create input files, and run simulations which include basic forward modeling, such as mixing samples, adding minerals or chemicals to a solution, etc. PHREEQC input files may contain samples from your AquaChem database or a Pure Water solutions. Before creating a PHREEQC Input file, it is recommended that you have some basic knowledge about PHREEQC. For more details, please refer to the PHREEQC User’s manual which is included with your AquaChem installation.

Option 7: PHREEQC (Advanced)

For a more advanced simulation, AquaChem provides links to the two versions of PHREEQC which are distributed by the USGS: PHREEQC-I and PHREEQC for Windows. These versions offer the full features of PHREEQC, including transport, inverse modeling, and kinetics. In order to do advanced modeling, and exploit all the features of PHREEQC, it is encouraged that you use one of these versions of PHREEQC.

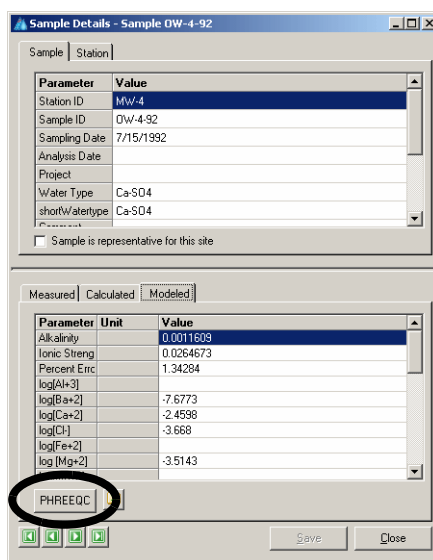
You may link either one of the mentioned programs to AquaChem in the **File > Preferences** and launch this program through the **Tools > Modeling > PHREEQC**

(Advanced) menu option. When this program is launched, an input file will be created automatically, and will contain any selected sample(s) from your AquaChem active samples list.

6.4.1 Calculate Saturation Indices and Activities

When you select this item from the **Tools > Modeling** menu, PHREEQC will run for each selected sample, and calculate saturation indices and activities for all modeled parameters which are defined in the current database structure. The results of the simulation will be automatically written back to the database for each selected sample, provided that the fields exist in the database. For example, to read back the calcite saturation index calculation, you need a field SI_Calcite specified in the **Database Options, Modeled Parameters** section.

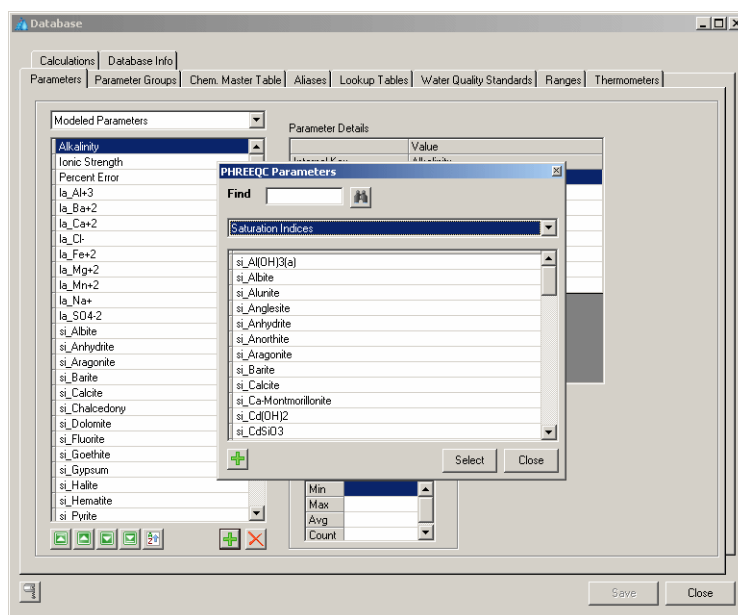
If you would like to calculate Saturation Indices (SI) or activities for **one** sample in your database, simply load the appropriate **Sample Details** window, then click on the **Modeled Parameters** tab. Once you have done this, click on the **PHREEQC** button, which is located in the lower left corner of this window (as circled below):



PHREEQC will then run in the background, and the modeled results will be saved automatically back to your database. There is no need to create and define input files.

If you would like to quickly calculate Saturation Indices (SI) or activities for **multiple** samples in your database, then you should use the **Calculate Sat. Indices and Activities** menu option.

To add additional modeled parameters to your database structure, you must load the Database options, under **File > Database > Parameters**. Then, select the **Modeled Parameters** from the combo box, and add the desired Saturation Indices and Activities to your database (as shown in the dialogue below).



An example of how to **Calculate Saturation Indices and Activities** is provided below:

Example

To model a sample from the **Demo.AQC** database:

- ☞ Select any sample from your Active Samples List.
- ☞ **Calculate Saturation Indices and Activities** from the **Tools > Modeling** menu.

You will then see a PHREEQC DOS window run in the background. Once this is finished, load the **Sample Details** window for this sample, then

- ☞ **Modeled Parameters** tab

In this window, you will see the modeled values for the available parameters. An example is shown in the figure below:

Parameter	Value
Station ID	MW-4
Sample ID	OW-4-92
Sampling Date	7/15/1992
Analysis Date	
Project	
Water Type	Ca-SO4
shortWwatertype	Ca-SO4

☐ Sample is representative for this site

Parameter	Unit	Value
Alkalinity		0.0011609
Ionic Streng		0.0264673
Percent Errc		1.34284
log(Al+3)		
log(Ba+2)		-7.6773
log(Ca+2)		-2.4598
log(Cl-)		-3.668
log(Fe+2)		
log (Mg+2)		-3.5143
lnv(Mn+2)		

PHREEQC

Save Close

You may now do further processing and analysis with these parameter values, such as plotting, reporting, and querying.

Note: In addition to using the **Tools > Modeling > Calculate Sat. Indices** option, you can also use the **Tools > Modeling > PHREEQC Basic** option to calculate saturation indices for multiple samples. From the PHREEQC Basic window, click on the Solutions branch of the tree (left side), then select the samples of interest and ADD them to your solutions list using the "Green Plus button". Then, click on the Simulation branch of the tree (left side), and click the Selected Output tab. Choose the equilibrium phases (minerals) you are interested in, then RUN the PHREEQC program. Once the run is finished, click on the **Spreadsheet Tab**, and you can find the **Saturation Indices**. These can be saved from PHREEQC basic, or simply Copied and Pasted into another program

6.4.2 Calculate pH

This tool can be used to estimate a missing pH value for an individual sample. The pH is calculated, based on the assumption that the groundwater comes from a carbonaceous aquifer and is in equilibrium with calcite. Usually a saturation index (SI) of 0 is thus specified. If a different SI is known to be typical for samples in the region, a different value can be specified.

Alternately, if a pH value was measured, and again the groundwater is assumed to be in close equilibrium with calcite, this tool can be used to validate the measured pH value. If the estimated theoretical pH is close to the measured pH, the pH measurement is probably true.

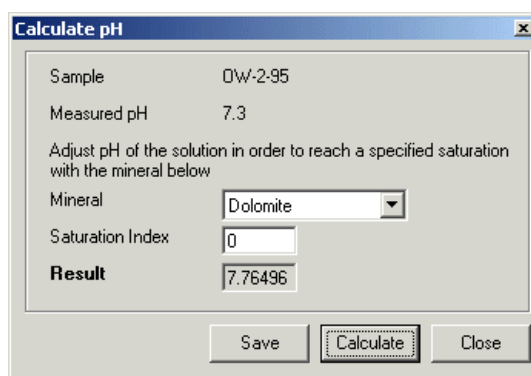
An example is provided below.

Example

To model the pH for a sample from the **Demo.AQC** database:

- ☞ Select any sample from your Active Samples List.
- ☞ **Calculate pH** from the **Tools > Modeling** menu.

You will then see the following dialogue:



The **Sample** and **Measured pH** will be filled in automatically.

- ☞ Select a **Mineral** from the combo box, usually calcite or dolomite.
- ☞ Enter a value for the **Saturation Index** for this Mineral. This may be from a previous simulation, or other known values.
- ☞ **[Calculate]**

You will then see a PHREEQC DOS window run in the background. Once this is finished, a calculated pH value will appear in the **Result** field.

To accept this calculated pH value,

- ☞ **[Save]**

Otherwise,

- ☞ **[Close]** to return to the main window.

6.4.3 Calculate Eh

In some cases, it may be very difficult to obtain meaningful Eh values by direct measurements. Several authors have recommended to calculate the Eh by means of the redox speciation with the Nernst Equation. Each Redox couple, e.g. $\text{Fe}^{2+}/\text{Fe}^{3+}$, $\text{Mn}^{2+}/\text{Mn}^{3+}$ gives an individual Eh value, which in cases of equilibrium conditions, should be

reasonable. The **Calculate Eh** menu item launches PHREEQC to search the selected sample for available redox couples and to calculate pe and Eh values for each couple. A sample of the dialog is shown below:

Calculate Eh

Sample MW-1-92

Measured Eh (mV)

Calculate the Eh based on the following redox couple

Redox couple	pe	Eh(mV)
Fe(2)/Fe(3)	5.3861	308
O(-2)/O(0)	14.3693	821

Close

The calculated Eh value may be compared to your measured Eh value.

In order to use this modeling utility, you must have data entered for a minimum of 1 redox couple (ex. Fe²⁺ / Fe³⁺), in the sample details window, **Measured** parameters tab.

6.4.4 Equilibrate with Minerals

This option allows you to evaporate a solution while staying in equilibrium with one or several minerals.

For additional information on this option please see “Modeling” on page 354 and “Equilibrate with Minerals” on page 174.

When this option is selected, the following dialog will load:

Equilibrate with Minerals

Sample Minerals Results

5 MW-1 7/28/1996 MW-1-96 Ca-Na-SO4 silty-gravel

Parameter	Unit	Value
Temperature	°C	14.2
Measured Alkalinity	mg/l CaCO3	
pH		7.3
Eh	mV	
Lithium	mg/L	
Potassium	mg/L	2
Sodium	mg/L	76
Calcium	mg/L	105
Magnesium	mg/L	26.2
Total iron	mg/L	0.02

Parameter Value

Evaporate (%) 20

Pump rate (m3/h) 15

Duration (days) 180

Calculate Close

This dialog consists of three tabs: **Sample**, **Minerals** and **Results**. Each is described below.

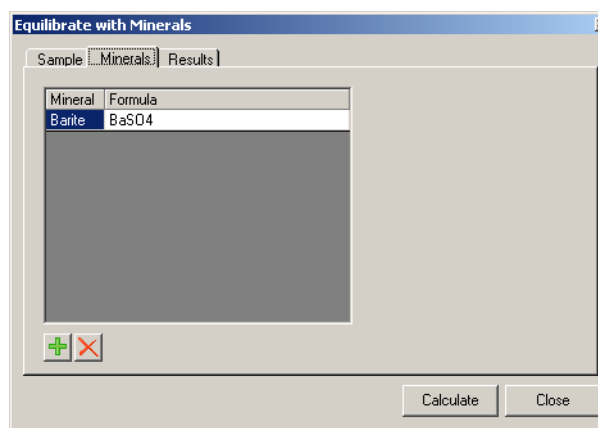
Sample

Sample information is shown in the top grid, i.e., station, sample date, sample_id etc. and its corresponding parameter information is shown below, in the left grid. In the adjacent right grid, you can enter values for **Evaporate (%)**, **Pumprate (m3/h)** and **Duration (days)**.

If the evaporation value is 0, the calculator will predict the mass of the precipitated minerals, based on the pumping rate and duration.

Minerals

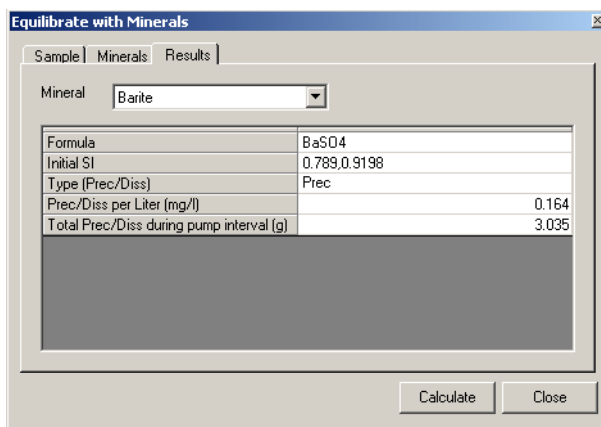
This tab allows you to specify the minerals to equilibrate the solution with.



Simply click the green button to add a new record in the minerals table. Click in the mineral field (highlighted in the image above) and then select the desired mineral from the combobox.

Click the **[Calculate]** button to generate the output values. These can be viewed in the Results tab.

Results

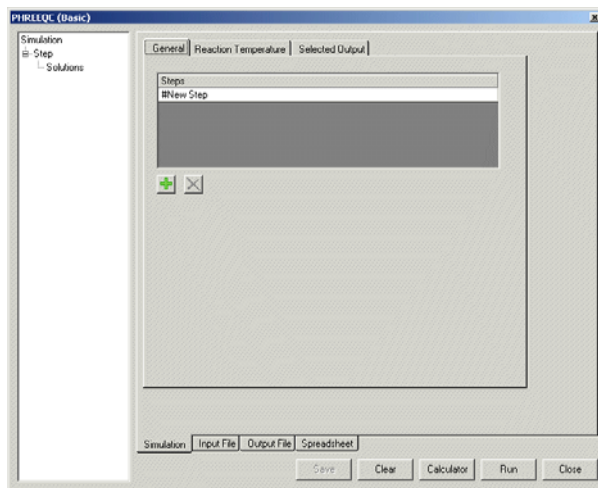


This tab displays the output results which include the chemistry of the remaining sample as well as the amount of mineral that was precipitated or dissolved (Prec/Diss). If a pumping rate and duration was specified in the Sample tab, the results are also expressed as total prec/diss during the pumping interval.

If you specified multiple minerals in the Mineral tab, you can recalculate the output with respect to different mineral by selecting the mineral from the Mineral combobox and clicking [**Calculate**].

6.4.5 PHREEQC (Basic)

This option loads the **PHREEQC (Basic) Input file** dialogue as shown below.



This graphical environment facilitates rapid development of simple PHREEQC simulations. Only the basic features of PHREEQC are supported; to take advantage of the more advanced features (Inverse Modeling, Transport, and Kinetics), you may utilize one of the USGS's PHREEQC versions, as explained below.

AquaChem provides direct access to your AquaChem project database samples for selecting solutions and building the input files. New solutions can be easily defined as needed, or existing solutions from previous simulations can be conveniently selected.

AquaChem also supports the use of flexible units for the various chemicals in solution.

Several simulations can be defined in one input file and processed in a single run. Reactions such as heating, evaporating, dissolving, or precipitating minerals can be conveniently simulated in step wise simulations, with all model parameters calculated for each step.

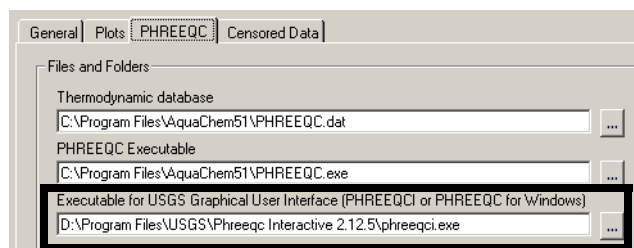
Once the PHREEQC input files have been prepared, the simulation can be launched directly from AquaChem. When the PHREEQC simulation is completed, you can copy the results from the spreadsheet view, and paste these into the Sample Details window for each individual sample. This will allow you to do further processing and analysis, such as plotting, reporting, and querying.

NOTE: Before using PHREEQC, you must ensure that you have defined the location of the PHREEQC.exe (executable), and PHREEQC.dat files. This can be done in the **PHREEQC Preferences** dialogue, available in the **File** menu.

For more details on designing a PHREEQC input file, please refer to Chapter 7: **Geochemical Modeling with PHREEQC (Basic)**.

6.4.6 PHREEQC (Advanced)

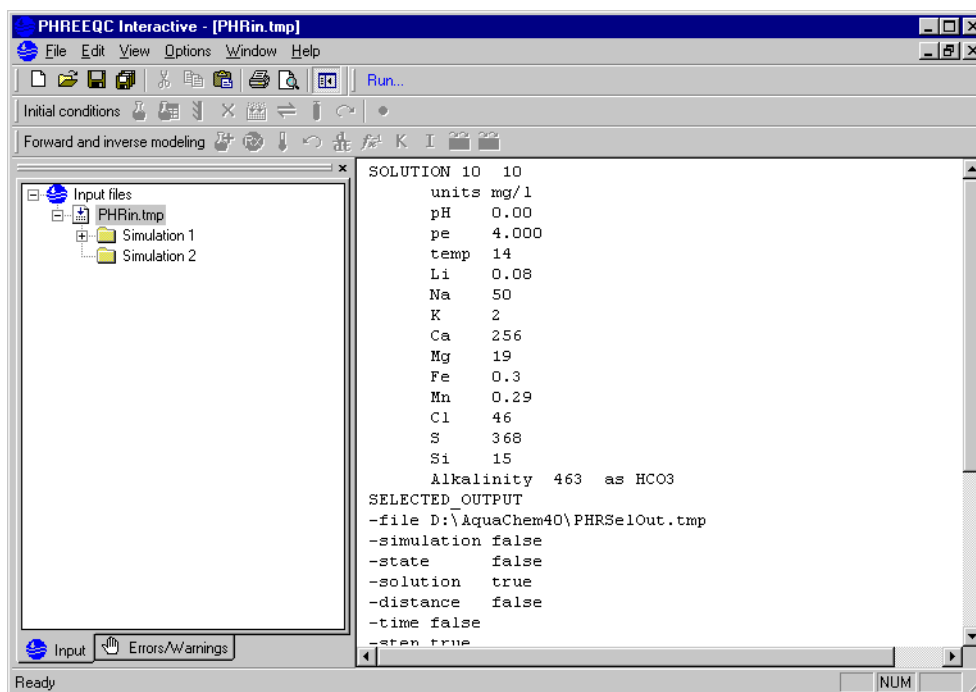
This option allows you to run advanced PHREEQC simulations, using either PHREEQC-Interactive or PHREEQC for Windows. The PHREEQC version that is loaded will be dependent upon the .exe file which is specified in the **Preferences** dialogue, as shown below:



In the third line under **Files and Folders**, you will see a field where you can specify the location of the PHREEQC executable file. If you are using PHREEQC-I, the file is called **phreeqci.exe**. If you are using PHREEQC for Windows, the file is called **phreeqc.exe**. Please point this to the appropriate folder on your system where the respective installation resides.

PHREEQC-Interactive

When you select the **PHREEQC (Advanced)** option from the **Tools > Modeling** menu, this will load the USGS's PHREEQC- Interactive Windows Interface, provided that this component was successfully installed during your AquaChem installation. An example of the PHREEQC-I window is shown below:



You may select individual or multiple samples in your active samples list. Any sample(s) which is (are) selected will be used as initial solution(s) when you load the PHREEQC-I program. You may then use the full features that PHREEQC-I has to offer. Once a PHREEQC-I simulation is completed, you must manually insert the results back into AquaChem by copy-and-paste, or by manually typing in the values.

NOTE: Before using PHREEQC-I, you must ensure that you have defined the location of the PHREEQCI.exe (executable). This can be done in the **Preferences>PHREEQC** dialogue, available in the **File** menu. If you did not install PHREEQC during your AquaChem installation, you may install this using the installation files located in the

PHREEQC folder of your AquaChem CD-ROM. The installation file is named: Phreeqc-2.12.1-669.exe.

For more details on PHREEQC-I, please refer to the user's manual "manual.pdf" which can be found in the 'Doc' folder, in your PHREEQC-I installation directory. Or, you may access the PHREEQC-I On-Line help, from within PHREEQC-I (in the Help menu).

PHREEQC for Windows

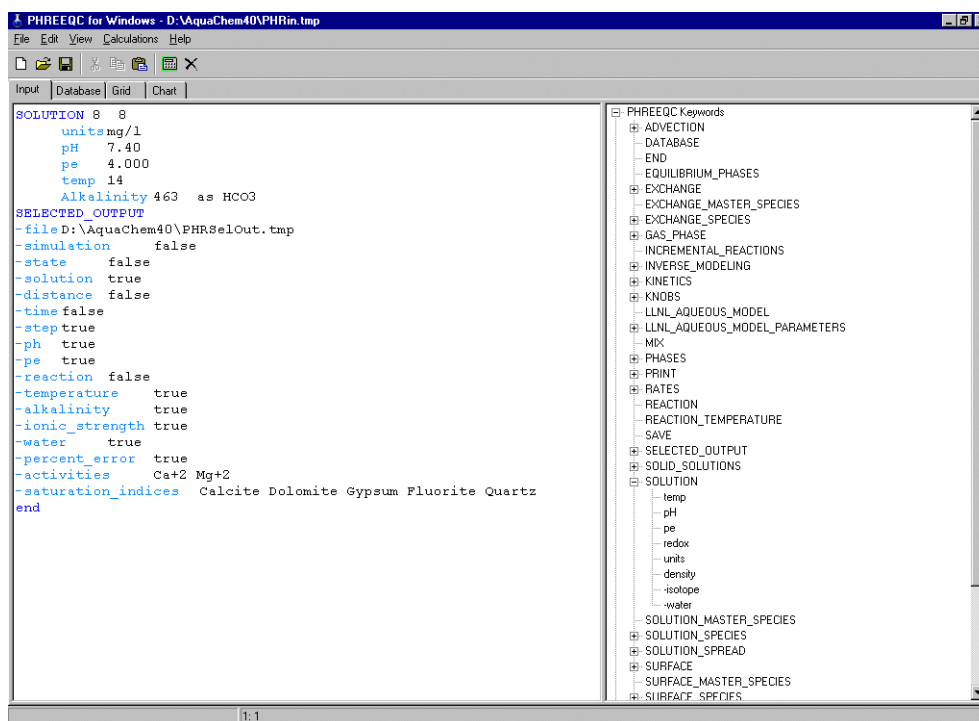
PHREEQC for Windows is a 32-bit Windows version of the geochemical model PHREEQC. PHREEQC for Windows contains the full functionality of PHREEQC v.2. The input files for the program are backward compatible with the normal version of PHREEQC. This means that you can use any file created with the normal version in PHREEQC for Windows. However, some options that are available in PHREEQC for Windows are not available in the standard version. The PHREEQC for Windows is developed by Vincent E.A. Post.

PHREEQC for Windows is not included with the AquaChem installation. If you would like to install this program, you may use the installation files which can be found on your AquaChem CD-ROM, in the PHREEQC folder. The installation file is named **psetup21204.exe**. PHREEQC for Windows is a public domain product, and may be downloaded from the web site below:

<http://www.geo.vu.nl/users/posv/phreeqc/index.html>

Once you have installed PHREEQC for Windows, you must define the location of the PHREEQC.exe (executable), in the **PHREEQC Preferences** dialogue, available in the **File** menu. Then, this executable will be launched when you select **PHREEQC (Advanced)** from the **Tools > Modeling** menu option.

An example of the input window for PHREEQC for Windows is shown below:



Similar to the PHREEQC-I, any sample(s) which is (are) selected in your active list will be used as initial solutions when you load the PHREEQC for Windows program. You may select individual or multiple samples in your active samples list. You may then use the full features that PHREEQC for Windows has to offer. Once a simulation is completed, you must manually insert the results back into AquaChem by copy-and-paste, or by manually typing in the values.

For more details on PHREEQC for Windows, please refer to the user's manual "ManualW.pdf", which can be found in your PHREEQC for Windows installation folder. Or, you may access the On-Line help, from within PHREEQC for Windows (in the Help menu).

6.4.7 Generate PHT3D Input

AquaChem v.5.1 can produce PHT3D input files for use with PHT3D in Visual MODFLOW versions 4.2 or higher.

Visual MODFLOW v.4.2 incorporates the reactive multi-component engine PHT3D. PHT3D couples the transport simulator MT3DMS and the geochemical package PHREEQC-2. Through the flexible, generic nature of PHREEQC-2, PHT3D can

handle a broad range of equilibrium and kinetic reactive processes, including aqueous complexation, mineral precipitation/dissolution and (multi-site) ion exchange. In the current version 4.2, Visual MODFLOW data entry is based on the same user interface used for the transport engines such as MT3DMS and RT3D. This has the following drawbacks: typically when using PHT3D, your model will be comprised of many more species than was the case with the former transport engines. Also, PHT3D only accepts special units, and solutions require special conditions such as perfect ion balance, total inorganic carbon entry instead of alkalinity, etc. Therefore laboratory results have to be transformed before the data may be used in PHT3D. To facilitate an efficient workflow, Visual MODFLOW includes an import feature, which allows the user to import solutions, minerals, gas composition and exchanger composition into a selected set of cells. AquaChem, in turn, allows for export of the information stored in its database into the format that can be imported into Visual MODFLOW.

Typically a PHT3D model is comprised of the following elements:

- initial composition of water in the aquifer
- composition of inflowing, infiltrating or injected water
- properties of the aquifer: mineralogical composition, amount and initial composition of exchanger places and gas composition

The operations described below are required to transform laboratory results into PHT3D ready input.

Solutions

All units must be expressed in mol/l, plus there must be a perfect ion balance between anions and cations. This is achieved by first running a PHREEQC simulation and allowing an automatic charge adjustment, typically on an element which is sufficiently abundant to achieve a balance and which is not considered important for the type of reactions studied through this model. In many cases chloride is a good candidate. Further, PHT3D requires total inorganic carbon as input as opposed to alkalinity which is usually reported in lab results. Again, this requires that the original solution is run and total inorganic carbon is read back from the simulation output file.

Minerals

Typically, the laboratory results are provided in ppm or percentage mineral related to 1kg of aquifer material. PHT3D requires the number of mols of mineral related to the mass of aquifer material in contact with 1 liter of pore water. This requires knowledge about the porosity and density of the aquifer material. For example, an aquifer having porosity of 20% and density of 2.7 g/cm^3 will have a volume of 4 liter of rock around every liter of pore water resulting in a mass of 10.8 kg of rock material. If this rock comprises 10% of calcite, the mols of calcite is calculated as $10.8 * 0.1 * \text{FMW}(\text{calcite}, 0.1 \text{ kg})$ 10.08 mols. This calculation needs to be repeated for each mineral.

Exchangers

PHT3D requires the amount of exchange places and the initial occupation of these places with various available ions, e.g. NaX, CaX₂, MgX₂, KX, etc. As in the previous case, the concentrations refer to 1 liter of pore water, though in reality, these values are rarely available. Values provided by the laboratory normally include the CEC (cation exchange capacity, which may be transformed into exchange places per liter of pore water in a similar way as discussed for the minerals. More often, the CEC itself has to be estimated based on the amount and type of clay minerals and presence of organic material. A formula that is often applied in this respect is given by Appelo and Postma, 1993:

$$\text{CEC} = 3.7 * \% \text{clay} + 3.5 * \% \text{ organic C}$$

If the type of the dominant clay material is known, then the CEC can also be estimated by multiplying the average CEC of this clay material with the percentage reported in the studied aquifer material.

All above methods only provide the places available for ion exchange, but do not indicate the initial occupation of these places. If an exchanger material assumed to be in equilibrium with a given solution, e.g. fresh water or sea water, then PHREEQC allows calculating this initial composition using the so-called implicit option for the exchange simulation. As for the solution, the PHREEQC output can then be used as input for PHT3D.

Create Input Files

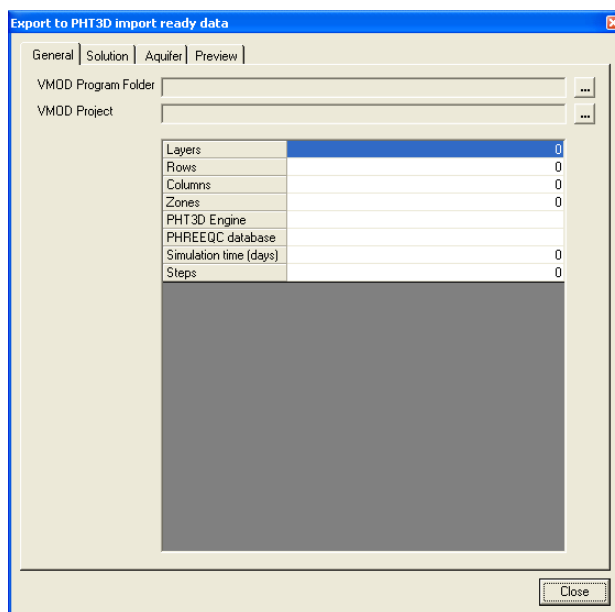
For models comprised of many species, data entry and associated conversion and transformation may turn out to be very tedious. AquaChem, therefore, includes an option to facilitate this data entry as explained below.

Requirements:

- A VMOD project must exist, since AquaChem reads information from the *.vmf file.
- A VMOD installation must exist on the local computer or network, since AquaChem needs to have access to the vmod.xml file in order to find the thermodynamic database.

To create a PHT3D input file:


- ☞ **Generate PHT3D Input** from the **Tools > Modeling** menu, and the screen depicted in figure below will appear.




Each tab is described in detail below:

General Tab

The **VMOD Program Folder** field allows you to select the VMOD installation folder (where the vmod.xml file is located).

Click the  button beside the VMOD Program Folder field, in the top right corner and navigate to the VMOD installation folder. Select the folder and click **[OK]**.

The **VMOD Project** field allows the user to specify the folders where the project .VMF file is located. AquaChem reads all required information from the XML file and displays it in the grid.

Click the  button beside the VMOD project field and navigate to the VMOD installation folder. Select the file and click **[OK]**.

The remaining tabs, *i.e.*, solution, aquifer and preview, provide an interface to define the composition that will be importable in PHT3D (solutions, minerals, gas composition, exchanger composition).

The first tab allows you to format the entry for one or several solutions. The solutions must have been previously entered in AquaChem using standard units. Another important precondition for assigning AquaChem concentrations to PHREEQC master species is that the master species are matched to AquaChem database parameters in the currently active thermodynamic database.

Solutions tab

The Solutions tab allows converting AquaChem solutions into PHT3D importable solutions including charge balance calculations and conversions to mol/L. When this tab is displayed, the grid below is shown. It lists all master species of the current thermodynamic database for the active VMOD project variant. The active compounds are marked in the third column by the internal concentration name, i.e., conc0001, conc0002, etc.

Element	M. Spec.	PHT3D	Conc.	Unit	mols/L
Temp					
pH					
pe					
H	H+				
H(0)	H2				
H(1)	H+				
E	e-				
O	H2O				
O(0)	O2	Conc014			
O(-2)	H2O				
Ca	Ca+2	Conc003		mg/L	
Mg	Mg+2	Conc008		mg/L	
Na	Na+	Conc013		mg/L	
K	K+	Conc007		mg/L	
Fe	Fe+2			mg/L	
Fe(2)	Fe+2	Conc005		mg/L	
Fe(3)	Fe+3	Conc006		mg/L	
Mn	Mn+2			mg/L	
Mn(2)	Mn+2				
Mn(3)	Mn+3				
Al	Al+3			ug/L	
Ba	Ba+2			mg/L	

Click the **[Select]** button to open the Station List dialog (shown below).

ID	Station	Date	Code	WATERTYPE	GEOLOGY	SYMBOL
1	Mw-1	8/15/1992	Mw-1-92	Ca-Na-SO4	silty-gravel	
2	Mw-1	6/1/1993	Mw-1-93	Ca-Na-SO4	silty-gravel	
3	Mw-1	6/15/1994	Mw-1-94	Ca-Na-SO4	silty-gravel	
4	Mw-1	7/30/1995	Mw-1-95	Ca-Na-SO4	silty-gravel	
5	Mw-1	7/28/1996	Mw-1-96	Ca-Na-SO4	silty-gravel	
6	Mw-1	6/15/1997	Mw-1-97	Ca-Na-SO4	silty-gravel	
7	Mw-1	8/1/1998	Mw-1-98	Na-SO4-Cl	silty-gravel	
8	Mw-3	8/8/1992	Mw-3-92	Ca-SO4-HCl	sandy-silt	
9	Mw-3	6/8/1993	Mw-3-93	Ca-HCO3-Si	sandy-silt	
10	Mw-3	6/15/1994	Mw-3-94	Ca-SO4-HCl	sandy-silt	
11	Mw-3	7/25/1995	Mw-3-95	Ca-HCO3-Si	sandy-silt	
12	Mw-3	8/2/1996	Mw-3-96	Ca-HCO3-Si	sandy-silt	
13	Mw-3	6/6/1997	Mw-3-97	Ca-HCO3-Si	sandy-silt	
14	Mw-3	7/30/1998	Mw-3-98	Ca-HCO3-Si	sandy-silt	

Select a sample from the station list, and click the **[Assign]** button and then **[Close]**.

All available concentrations from the selected sample will be copied to the “Conc” column in their original format.

The **Charge Balance** combo box allows you to select an element that will be used for calculating the charge balance. Please note that only the active PHT3D components will be used for the PHREEQC simulation. Therefore an AquaChem sample, that is perfectly balanced, but includes species which are not active in PHT3D, still needs to be balanced.

All available active PHT3D species will be written to a PHREEQC input file and the charge balance is calculated as well as redox species, e.g., Fe(2), Fe(3) from total Fe. All results are read back in mols/L column in the grid immediately (shown below). You may edit the concentration column manually however you will need to click the **[Run]** button to launch the PHREEQC simulation manually. When changing the element used to calculate the ion balance, the simulation is launched automatically.

Element	M. Spec.	PHT3D	Conc.	Unit	mols/L
Temp					14.8
pH					7.15
pe					4
H	H+				
H(0)	H2				
H(1)	H+				
E	e-				
O	H2O				
O(0)	O2	Conc014			0.0000E+00
O(2)	H2O				
Ca	Ca+2	Conc003	125	mg/L	5.8380E-03
Mg	Mg+2	Conc008	22	mg/L	9.0584E-04
Na	Na+	Conc013	80	mg/L	3.4834E-03
K	K+	Conc007	1.5	mg/L	3.8401E-05
Fe	Fe+2		0.96	mg/L	
Fe(2)	Fe+2	Conc005		mg/L	1.5903E-05
Fe(3)	Fe+3	Conc006		mg/L	1.3041E-06
Mn	Mn+2		0.6	mg/L	
Mn(2)	Mn+2				
Mn(3)	Mn+3				
Al	Al+3			ug/L	
Ba	Ba+2		4	mg/L	

Use the format combo box to specify the appropriate format for the generated input file. You can choose from **initial concentration** or **recharge concentration**.

For more information on recharge concentration format, see “Preview Tab” on page 375

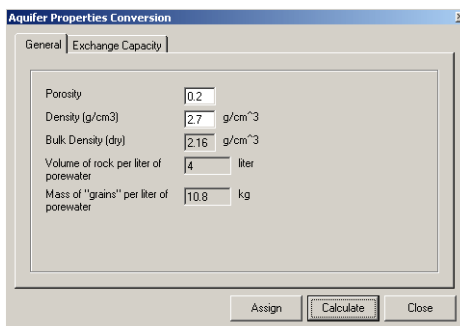
Aquifer tab

The Aquifer tab allows you to define the mineral composition of the aquifer. Note that PHT3D uses concentrations per liter of bulk aquifer material rather than per liter of porewater used by PHREEQC.

First, define the bulk density by entering the value in the **Bulk Density** text field. This value can be calculated based on the porosity and the density of grains. Most rock

forming minerals have a density of 2.6-2.75 g/cc. Porosity can vary more widely from a few percent up to 60% for clays.

Click the  button to access the AquaChem bulk density calculator (shown below).



The dialog box is titled "AquaChem Properties Conversion". It has two tabs: "General" and "Exchange Capacity". The "General" tab is active. It contains several input fields with their corresponding units and calculated values:

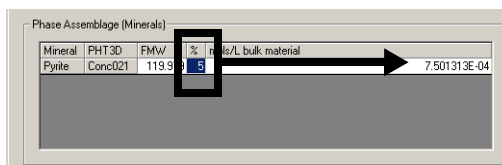
Property	Value	Unit
Porosity	0.2	
Density (g/cm ³)	2.7	g/cm ³
Bulk Density (dry)	2.18	g/cm ³
Volume of rock per liter of porewater	4	liter
Mass of "grains" per liter of porewater	10.8	kg

At the bottom right, there are three buttons: "Assign", "Calculate", and "Close".

Enter a **Porosity** and **Density** value and then click the **[Calculate]** button to generate the Bulk Density value. Click **[Close]** and enter the generated value into the **Bulk Density** text field in the Aquifer tab.

Minerals

In the **Phase Assembly (Minerals)** frame, enter the estimated or measured percentages for **each** listed mineral in the **%** column. The respective mols/L of aquifer bulk material will be calculated in the last column (shown below).




The table is titled "Phase Assembly (Minerals)". It has five columns: Mineral, PHT3D, FMW, %, and mols/L bulk material. The first row is for "Pyrite" with PHT3D "Conc021" and FMW "119.9". The percentage value "5" is entered in the % column and is highlighted with a black box. An arrow points from this box to the "mols/L bulk material" column, which contains the value "7.501313E-04".

Mineral	PHT3D	FMW	%	mols/L bulk material
Pyrite	Conc021	119.9	5	7.501313E-04

Exchanger Sites

The definition of an exchanger assemblage requires two basic inputs: the amount of total available sites and the initial distribution of these sites among the active PHT3D exchanger components. Because these values are not always directly measured, you can use the AquaChem calculator to generate estimated values.

The first step of calculating an exchanger composition consists of estimating the amount of exchanger sites in the aquifer material that is in contact with 1 liter for porewater. Click the  button and the screen shown below will be shown.

Under the **General** tab, calculate the Bulk Density by entering grain density and porosity values or simply enter the bulk density if the value is already known.

Next click on the **Exchange Capacity** tab.

For this example the exchange places are to be estimated based on a known percentage of 10% clay and 5% organic matter. Enter these percentage values in the first two text boxes. Click the **CEC estimated** radio button. Once the **[Calculate]** button is clicked, the CEC in meq/100g soil will be calculated and multiplied by the mass of grains calculated on the first tab.

Copy the calculated value to the clipboard (highlight value, right-click and select copy) and then paste the value into the **Exchanger sites** field of the **Minerals** tab.

There are two ways of entering the distribution of sites among Exchanger species: **Explicit** requires that you distribute the sites manually among the species, based on laboratory results or estimation. The **Implicit** method allows selecting a solution,

equilibrates the exchanger assemblage with this solution, and then reads the results back to the grid. Both methods are described in more detail in the following sections.

Explicit

If the distribution of specific exchange places (CaX2, NaX, KX, etc.) is known, enter the percent values manually in the grid below for each exchanger.

Exchanger

Exchanger sites: 1.82000 mol/L bulk material

☒ Explicit ☐ Implicit: exchanger is in equilibrium with sample:

Exchanger	PHT30	%	mol/L bulk material
Na+	Conc00	44	4.448e-002
Ca+2	Conc01	51	7.554e-001
Mg+2	Conc01	18	1.304e-001
SO4-2	Conc01	0	0
Ba+2	Conc01	01	1.677e-004

Implicit

Alternatively you may assume that the exchanger is in equilibrium with a solution. Typically for exchange reaction simulations, the aquifer including its exchangers is in equilibrium with an initial solution. This system is then flushed with a solution of a different composition (e.g. saltwater if the initial solution was freshwater or vice versa). The first solution can be used to calculate the distribution of individual exchange species (CaX2, NaX, KX, etc.).

Select the **Implicit** option, then press the button and select the required sample from the Station List dialog (shown below).

Station List

Find: in <all>

ID	Station	Date	Code	WATERTYPE	GEOLOGY	SYMBOL
1	Mw-1	8/15/1992	Mw-1-92	Ca-Na-SO4	silty-gravel	
2	Mw-1	6/1/1993	Mw-1-93	Ca-Na-SO4	silty-gravel	
3	Mw-1	6/15/1994	Mw-1-94	Ca-Na-SO4	silty-gravel	
4	Mw-1	7/30/1995	Mw-1-95	Ca-Na-SO4	silty-gravel	
5	Mw-1	7/28/1996	Mw-1-96	Ca-Na-SO4	silty-gravel	
6	Mw-1	6/15/1997	Mw-1-97	Ca-Na-SO4	silty-gravel	
7	OW-2	8/1/1998	Mw-1-98	Na-SO4-Cl	silty-sand	
8	OW-2	8/8/1992	Mw-3-92	Ca-SO4-HCl	silty-sand	
9	OW-2	6/8/1993	Mw-3-93	Ca-HCO3-Si	silty-sand	
10	OW-2	6/15/1994	Mw-3-94	Ca-SO4-HCl	silty-sand	
11	OW-2	7/25/1995	Mw-3-95	Ca-HCO3-Si	silty-sand	
12	OW-2	8/2/1996	Mw-3-96	Ca-HCO3-Si	silty-sand	
13	OW-2	6/6/1997	Mw-3-97	Ca-HCO3-Si	silty-sand	
14	OW-2	7/30/1998	Mw-3-98	Ca-HCO3-Si	silty-sand	

Assign Close

Once selected, click the **[Assign]** button.

This will automatically generate a PHREEQC simulation, equilibrating an exchanger having the specified number of exchange places with this solution. The calculated

exchange species are then extracted from the PHREEQC output file and inserted into the grid.

Exchanger

Exchanger sites: 1.82000 mol/L bulk material

☐ Explicit ☒ Implicit, exchanger is in equilibrium with sample:

Exchanger	PHT3D	%	mol/L bulk material
NaX	Conc009	66	4.839e-002
CaX2	Conc010	64	7.396e-001
MgX2	Conc011	42	1.350e-001
SiX2	Conc012	0	0
BaX2	Conc013	36	6.641e-003

Preview Tab

The preview tab provides a summary of the prepared data for PHT3D import. Three different entries are created: initial concentration, mineral composition and exchanger composition. Each grid contains a **[Save]** button which allows you to export the grid data to an importable text file (indicated in the image below).

Export to PHT3D import ready data

General | Solution | Aquifer | Preview

Species

C(4)	Cl(4)	Ca	Cl	Fe(2)	Fe(3)	K	Mg
0.0023729	0	0.0031203	0.0094831	0	0	0	0.0009053

Save Species

Minerals

Calcite	Dolomite	Anhydrite
9.92153E-04	9.52263E-04	3.22153E-04

Save Minerals

Exchangers

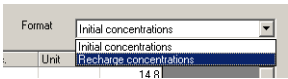
NaX	CaX2	MgX2	SiX2	BaX2
0.03796	0.741	0.1473	0	0.001227

Save Exchangers

Close

Exchanger and mineral composition can only be defined under the *initial concentrations* format. The respective datafiles are time independent. Solutions may also be bound to time dependant boundary conditions such as recharge and point sources. Among those, only the creation of recharge concentrations is supported for the time being.

To create recharge concentrations, simply select **Recharge concentrations** from the **Format** combo box in the **General** tab.



The preview grid for solutions will now be filled with multiple records (shown below). The number of time steps and the simulation duration is read directly from the project.vmf file and is also displayed in the project setting grid on the General tab. For the time being, the composition of every step is kept constant. This grid is editable; you may add or delete rows and change the start/stop values for every row. For example, if the solution includes a parameter for temperature, you can manually change this value for every time step, while the composition of the water remains constant.

Start Time (c)	Stop Time (c)	Ca	Mg	Na	Cl	Calcite
0	365	0.00337	0.00107	0.0029593	0.0092387	0.0028185
365	730	0.00337	0.00107	0.0029593	0.0092387	0.0028185
730	1095	0.00337	0.00107	0.0029593	0.0092387	0.0028185
1095	1460	0.00337	0.00107	0.0029593	0.0092387	0.0028185
1460	1825	0.00337	0.00107	0.0029593	0.0092387	0.0028185

Once the grid data has been saved in imported text files, they can be imported into VMOD/PHT3D.

7

Geochemical Modeling with PHREEQC (Basic)

This chapter presents an overview of the modeling capabilities of PHREEQC and an introduction on how to create PHREEQC Input files using the basic version of PHREEQC included within the AquaChem interface.

Here you will find an introduction to the AquaChem Interface to PHREEQC, PHREEQC Window Layout, Creating PHREEQC Input Files, and Running PHREEQC Simulation.

AquaChem uses the batch version of PHREEQC, which allows to integrate this model seamlessly into AquaChem. For details on the Graphical User Interfaces PHREEQC-I, or PHREEQC for Windows, please refer to the User's Manuals included with these programs. Or, for a basic description, please refer to PHREEQC (Basic) section in Chapter 6.

7.1 AquaChem Interface to PHREEQC

PHREEQC is a computer program for speciation, batch-reaction, one-dimensional transport, inverse geochemical calculations, and much more. For more than twenty years, the USGS's PHREEQC has been the proven standard for aqueous geochemical modeling. PHREEQC is derived from the FORTRAN program PHREEQE. As the name implies, the original FORTRAN code has been entirely rewritten in the C programming language and this new version has eliminated many of the deficiencies and limitations of earlier version of PHREEQE, while adding many new powerful analytical capabilities.

In the past years the authors of PHREEQC have not only added many more powerful features to the program, they have also implemented two very easy-to-use interfaces for writing input files and analyzing output files for the PHREEQC program. The respective programs PHREEQC-I and PHREEQC for Windows are included with your AquaChem installation CD; you may link your AquaChem samples in your database to either one of these programs, provided that these programs are installed, and linked in the **File > Preferences** dialogue. These interfaces may be called from within

AquaChem, and this will allow you to run advanced PHREEQC simulations using your AquaChem samples. For more details, please refer to Chapter 6.

The **PHREEQC (Basic)** option provides a GUI to the basic features of PHREEQC. You may run simple simulations based on analysis stored in the AquaChem database; this option also allows you to create PHREEQC input files, can be later opened with more sophisticated graphical user interfaces such as PHREEQCI or PHREEQC for Windows versions of PHREEQC.

Although this chapter describes how to create PHREEQC input files with the PHREEQC (Basic) version, the concept of hydrochemical modeling and the PHREEQC program will not be discussed in detail. It is recommended that you read the PHREEQC user's manual before performing any complex hydrochemical simulations. For your reference, the PHREEQC user's manual and source code may be downloaded from the following web site:

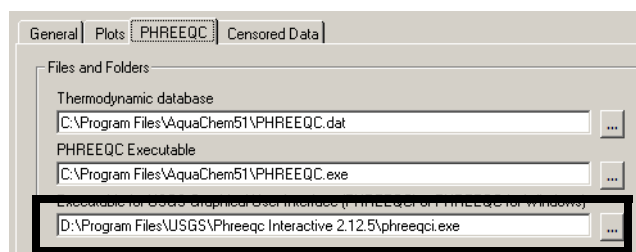
http://wwwbrr.cr.usgs.gov/projects/GWC_coupled/phreeqc/.

In addition, the PHREEQC user's manual is available on the AquaChem installation CD-ROM, and in your AquaChem installation folder.

7.1.1 Preferences for PHREEQC


The instructions in this chapter will be based on the assumption that you are using the PHREEQC executable file (PHREEQC.exe - PHREEQC version 2.8) included with AquaChem. By default, the PHREEQC batch executable file and the PHREEQC thermodynamic database files are automatically installed in the default folder, C:\Program Files\AquaChem. If you have installed AquaChem in a directory other than the default installation directory, or if aquachem is not yet linked to PHREEQC, the **Tools/Modeling/PHREEQC (Advanced)** option will be inactive (greyed out) on your computer, and you will need to manually set up the link between AquaChem and the PHREEQC thermodynamic database and executable files.

To do so, select **File** from the main menu and then select **Preferences**. The following **Preferences** dialogue will appear. In this dialogue, select the **PHREEQC** tab.



Under the **Files and Folders** frame, press  button beside the **Thermodynamic Database** field and select the path and filename from the **PHREEQC Thermodynamic**

Database dialogue. From this dialogue, select either **phreeqc.dat**, **Wateq4f.dat** or **Minteq.dat**. In addition, in the **PHREEQC Executable** field select the path and filename for the **phreeqc.exe** file. Finally, specify the path and filename for the USGS PHREEQC Executable file (which is used in the PHREEQC (Advanced) option).

If the default path and filename are incorrect, you can browse the directory by clicking on the  button and select the filename.

By default, all PHREEQC input and output files will be saved in your AquaChem installation folder (default is C:\Program Files\AquaChem).

Once you are finished, click **[Save]** then **[Close]**. You are now ready to run PHREEQC.

7.1.2 The PHREEQC Thermodynamic Database Link

One of the biggest advantages of the AquaChem PHREEQC interface is the direct link between the two databases of these programs. The link allows you to use samples from the AquaChem database and read them into the PHREEQC input data file.

In order to let AquaChem know which database parameter matches which PHREEQC element, some minor modifications have been made to each of the thermodynamic databases included with AquaChem. The link between AquaChem and the thermodynamic databases is established by adding the AquaChem parameter name as a comment for each matching parameter in the Masterspecies section of the database file. Each comment is preceded by a '#' character to flag the AquaChem parameter descriptor.

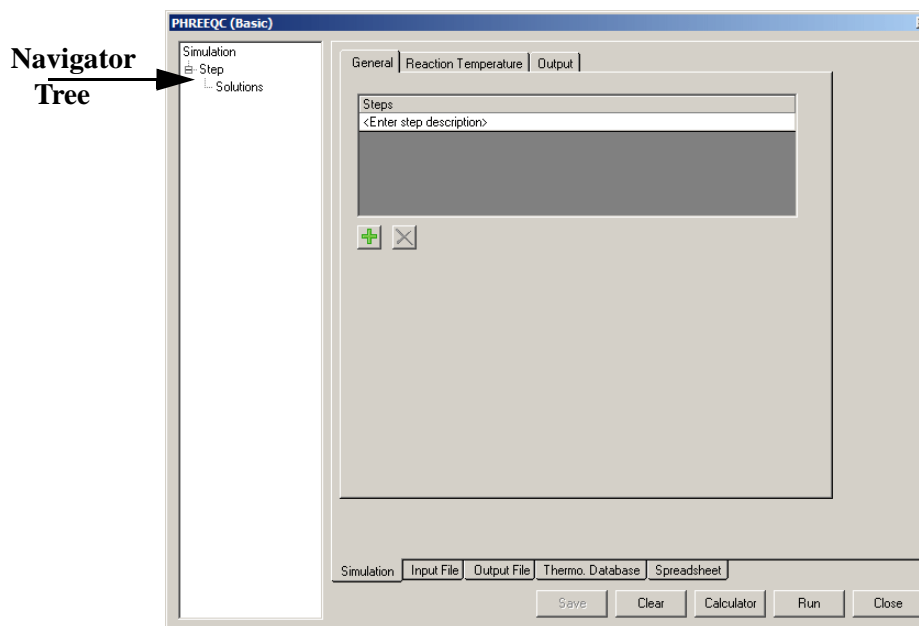
For example, the following lines are taken from the Masterspecies section of the **phreeqc.dat** file:

Ca	Ca+2	0.0	Ca	40.08	#CA
Mg	Mg+2	0.0	Mg	24.312	#MG
Na	Na+	0.0	Na	22.9898	#NA
K	K+	0.0	K	39.102	#K
Fe	Fe+2	0.0	Fe	55.847	#FE
Fe(+2)	Fe+2	0.0	Fe		
Fe(+3)	Fe+3	-2.0	Fe		
Mn	Mn+2	0.0	Mn	54.938	#MN

These modifications have already been defined for each of the thermodynamic database files provided with AquaChem. You have to redefine this only if you change the thermodynamic database.

7.2 PHREEQC Window Layout

To access the PHREEQC input dialogue, select **Tools** from the main menu of AquaChem and **Modeling** and then **PHREEQC (Basic)**. This will load the **PHREEQC (Basic) Input** dialogue as shown in the following figure.



The **PHREEQC Input** dialogue consists of several options and settings that apply to the entire PHREEQC simulation.

The left side of the dialogue includes a **Navigator Tree**, allowing you to quickly add and modify the reaction options for the current simulation.

Along the lower part of the **PHREEQC Input** dialogue there are four tabs:

- | | |
|--------------------|---|
| Simulation | This allows you to define the simulation options (main input dialogue). |
| Input File | This provides a view of the PHREEQC input file, in text format. |
| Output File | This provides a view of the complete PHREEQC output file, in text format. |

Spreadsheet This provides a spreadsheet preview of selected parameters from the PHREEQC output file in a spreadsheet format.

The bottom of the **PHREEQC Input** dialogue contains several buttons:

Save The save button allows you to save the contents of the input or output file. If the **Input File** tab is selected, the **[Save]** button will allow you to save the input file as a **.PQI** file. If desired, this file may be loaded into PHREEQC-I, and the simulation may be run using this version. If the **Output File** tab is selected, the Save button will allow you to save the output file as a **.PQO** file.

Clear The **[Clear]** button clears the contents of the input file, and allows you to create a clean, new input file.

Calculator The **[Calculator]** button loads the **Aquifer Properties Conversion** dialogue and allows you to calculate and convert various aquifer properties you require for some of the PHREEQC simulations.

Run The **[Run]** button runs PHREEQC using the current simulation options. This button loads the PHREEQC input file and executes the PHREEQC simulation. The **[Run]** button should only be used after you have specified all of the required run options and you are ready to start the PHREEQC simulation. It is recommended that you verify the input file before running the simulation.

Close The **[Close]** button closes the PHREEQC input dialogue.

There are four tabs associated with the **[Calculator]** button: **General**, **Surface Complexation**, **Exchange Capacity**, and **Mineral Assemblage**. These are shown and described below:

The screenshot shows a dialog box titled "Aquifer Properties Conversion" with a standard Windows window border. It has four tabs: "General", "Surface Complexation", "Exchange Capacity", and "Mineral Assemblage". The "General" tab is currently selected. Inside the dialog, there are four input fields with labels to their left: "Porosity" with a value of "0.2", "Density (g/cm3)" with a value of "2.7", "Volume of rock (liter) per liter of pore water" with a value of "0" and the unit "liter" to the right, and "Mass of grains (kg) per liter of pore water" with a value of "0" and the unit "kg" to the right. At the bottom right of the dialog, there are two buttons: "Calculate" and "Close".

General

Since PHREEQC always assumes a solution volume of 1.0 liters, the **General** calculator allows you to easily calculate the **Volume of rock (liters) per liter of porewater** and **Mass of grains (kg) per liter of porewater** for the aquifer based on a solution containing 1.0 liters of porewater. Simply enter the input parameters (**Porosity**, **Density**), and press the **[Calculate]** button. This provides the necessary values for all calculations shown below.

Surface Complexation

The moles of surface places (**sites**) depends on the content of hydrous ferric oxides in the aquifer (**Sites/mol ferrous iron**). **Fe** is generally a measured parameter but you must estimate the percentage of iron (**Weight % Fe**) that is in the form of hydrous iron (**% hydrous ferric oxides of total iron**). Since PHREEQC always asks for mols, you also need a formula weight value for your iron minerals. Hydrous iron is a mixture of several minerals, so a suggested **Gram formula weight for oxides** is **89** which is used as a default value.

The screenshot shows a software window titled "Aquifer Properties Conversion" with four tabs: "General", "Surface Complexation", "Exchange Capacity", and "Mineral Assemblage". The "Surface Complexation" tab is active. It contains a table with the following fields and values:

Parameter	Value
Weight % Fe	2
% hydrous ferric oxides of total iron	10
Gram formula weight for oxides:	89
Sites/mol ferrous iron (mol)	0.2
Surface sites /l Porewater	0
Mass of ferrous iron (g)	0

Exchange Capacity

The Cation Exchange Capacity (CEC) is normally expressed as meq/100g of soil. PHREEQC however uses meq/l of porewater. There are several ways of estimating the number of exchangeable sites and express it in the unit used by PHREEQC.

If you know the CEC value you may directly enter it and calculate the exchange places (**Sites**) within the given aquifer volume. Otherwise you need to estimate the CEC. The first option lets you select a clay mineral from a list and will use an average CEC value. Choose this option if you know the dominant clay mineral of your aquifer. If you know only the percentage but not the mineralogy of your clay minerals you may use a formula derived by Appelo (1993) to estimate the CEC:

$$\text{CEC} = 0.7 * \text{clay\%} + 3.5 * \text{org material\%}.$$

Mineral Assemblage

The **Mineral Assemblage** option lets you convert an analysis usually expressed as weight percent to mol of mineral per liter of porewater. Simply select a mineral from the combo box, and specify a **Weight %**. Press **[Calculate]** and the amount of mols will be calculated.

Mineral	FMW [g]	Weight%	mol
Calcite	100.0872	10	10.791
Dolomite	184.4014	5	2.928

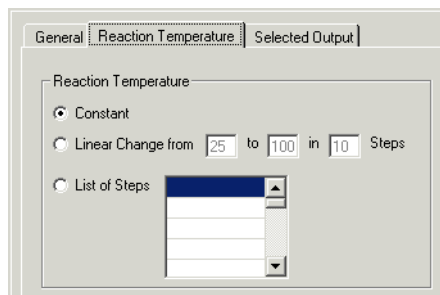
There are three tabs associated with the **Simulation** tab which are shown and described below:

General

The **General** tab contains a summary of the steps defined for the current simulation.

Reaction Temperature

This tab provides options to specify the reaction temperature, or step-wise changes in the reaction temperature. The **Reaction Temperature** options can be seen in the dialogue shown below.



The **Reaction Temperature** tab provides options for specifying the desired temperature settings during a modeled reaction. Normally the temperature of the selected sample stays **Constant** (default setting) during a reaction and it will use the solution temperature as entered in the **Solutions** option selected from **Navigator tree**. However, PHREEQC provides you the options to change the temperature linearly (**Linear change from...to...in...Steps**) from a starting temperature to an ending temperature in a specified number of steps or in steps during the course of a simulation. The **List of Steps** option allows you to specify the exact temperature at specific steps during the reaction and the temperatures will be applied in the order they are listed.

Output

The **Output** tab allows you to access the various output details to include in the standard PHREEQC results output file (**.PQO** file extension) or to save simulations directly back to the database. PHREEQC will generate an output file every time you run a simulation. However, this file is very detailed and may contain more information than you need to solve a particular problem. Its format is also not suited for plotting or presenting simulated data. This tab lets you extract individual parameters from the detailed output file and write them into a spreadsheet.

The options available under the **Selected Output** dialogue are shown in the figure below:

The screenshot shows the 'Output' tab of the PHREEQC interface. The 'Selected Output' section is active, displaying six columns of checkboxes for selecting data to be outputted:

- Totals:** ☐ H, ☐ H(0), ☐ H(1)
- Activities:** ☐ H+, ☐ e-, ☐ H2O
- Molalities:** ☐ H+, ☐ e-, ☐ H2O
- Saturation indices:** ☐ Calcite, ☐ Aragonite, ☐ Dolomite
- Equilibrium phases:** ☐ Calcite, ☐ Aragonite, ☐ Dolomite
- Gas phases:** ☐ CO2(g), ☐ O2(g), ☐ H2(g)

Below these columns, there is a radio button for 'Prepare simulated solution output for storage in database'. If selected, it includes a dropdown menu labeled 'Assign simulated value to Station:'.

Selected Output

With the selected output option, you may extract specified activities, saturation indices, gas phases, etc. into a spreadsheet type file.

Simply choose the parameters you want to save to the tabular output file. By default the parameters **pH**, **pe**, **Temperature** and **Ionic Strength** are included in the output.

Totals Total amount of a masterspecies, this value should be equal to the entered concentrations


Activities Amount of aqueous species that is available for reactions.

Molalities Concentrations of aqueous species

Saturation indices Saturation state of minerals

Equilibrium phases: Amount of minerals present in the system

Gas phases: Amount of gas phases present in the system

When you have successfully run the simulation, AquaChem allows you to save the output file. To do so, press the  button, and enter a filename.

Prepare simulated output for storage in database

If the database is selected as the target output, then the simulated results for created samples will be saved as new sample records. This option is only applicable if a simulation creates a new solution, e.g. when mixing several existing samples, when equilibrating an existing sample with minerals, or when adding chemicals. Since every sample needs to have an assigned station, you should create one or several dummy simulation stations, to which you may assign your simulated samples in order to easily distinguish them from real samples. If the sample comment field exists and is identified and mapped in the **File>Database>Aliases** section, then the type of simulation will be written automatically to the sample.

Simulated samples do not necessarily belong to a station, e.g. when saving the mixture between 2 samples, the saved simulated composition would belong to no specific station. However, AquaChem needs a station to be assigned to every sample, therefore the **Save simulated solution output to database** option requires you to define the station for the new samples to be created. You may create a dummy station names, such as “PHREEQC Simulation” that may be used for this purpose.

Save Options


The PHREEQC input and output files are saved by Aquachem under default filenames which will be overwritten with every new PHREEQC run. Using the “**Save Solution as**” option in the step frames, allows you to save the current simulation under a defined name. You may afterwards use this file for your documentation or continue to model with one of the USGS GUIs that provide more advanced coverage of the PHREEQC features.

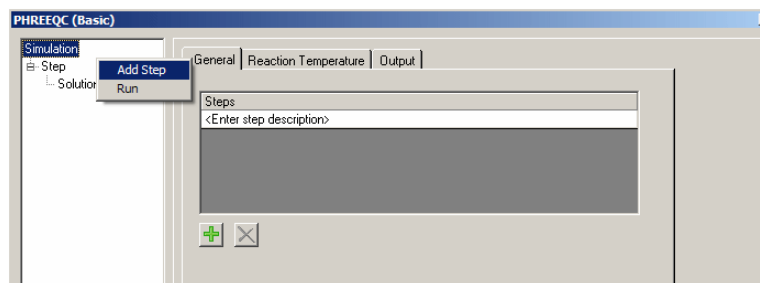
7.3 Creating PHREEQC Input Files



7.3.1 Simulation - Steps

A PHREEQC simulation may consist of several steps (e.g you may define and mix 2 samples in the first step and equilibrate the mixing result in a second step).

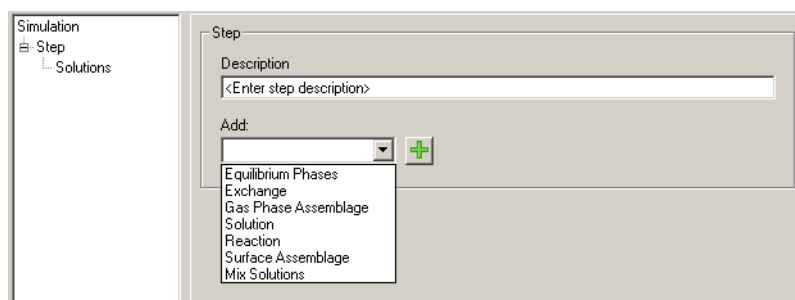
Under the **General** tab, you will see a list of **Steps** (if multiple steps have been defined).

In order to add a simulation step you may right click on **Simulation** - the top item of the **Navigation tree** - and select the **Add Step**. Or click the  button below the **Steps** frame.



You can enter a name for the **Step** in the **Description** field, and choose the step properties. Click  button under the **Add** and choose an item from the list, and press the  button.

After a new step has been added, double click on it or click on the Step in the **Navigator tree** to produce the Step description dialogue:



This new item will now appear in the **Navigator tree** on the left side of your display. To activate the properties for this item, expand the **Navigator tree** and click once on this item.

To modify the **Step** properties, you have two options:

- [1] Double click on the **Step** name from the main dialogue, OR
- [2] Select "**Step**" from the **Navigator tree** on the left side of your display.

Options for the reaction steps can also be created and modified as needed. These include:

- **Add Initial Conditions** and
- **Forward Modeling**

These options are described in detail in the following sections.

Add Initial Conditions

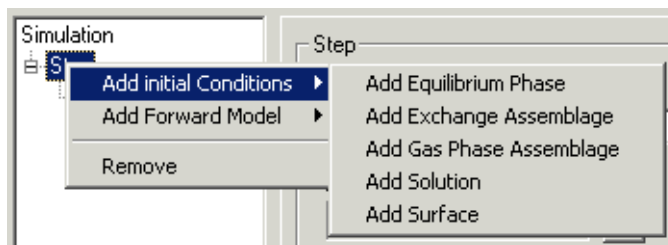
Add Equilibrium Phases

Add Exchange Assemblage

Add Gas Phase Assemblage

Add Solutions

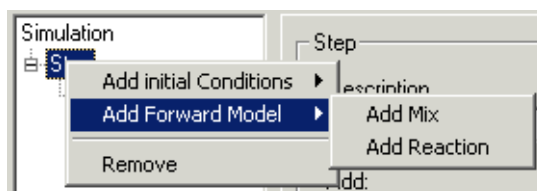
Add Surface Assemblage



Forward Modeling

Add Mix Solutions

Add Reactions



7.3.2 Equilibrium Phases

Adding Equilibrium mineral phases is similar to adding reactions. With the **Add Reaction** options (described below) you add a specified amount of a phase, but with the **Add Equilibrium Phase** options the amount of mineral phase added is limited by a specified saturation index.

To specify an equilibration of your solution with Mineral Phases, right click on the **Steps** options and select **Add Initial Conditions** from the pop-up menu and then select **Add Equilibrium Phases**.

Equilibrium Assemblage

Each mineral phase assemblage is identified by a unique **Phase** assemblage **Number** and a **Description**. You may enter multiple phases by specifying a phase number from # to # (ex. Number 1 to 2). The assemblage number is only important if you intend to use the input file later for transport modeling using PHREEQCI or PHREEQC for Windows.

- Select **Equilibrium Phases1** and an **Equilibrium Phases** dialogue will appear as shown below.

Simulation
└─ Step
 └─ Solutions
 └─ Equilibrium Phases1

Equilibrium Phases

Description: <Enter Equilibrium Phase Assemblage description>

Number: 1 to

Phase	SI	Mols	Alt. phase	Mols
Calcite		0		

+ X

Equilibrate Phase Assemblage with solution: (If not specified, first solution in simulation will be used) ☐

Save Solution as

For each mineral phase you choose, you can specify the following properties:

SI: This is the Saturation Index of the selected mineral phase (degree of saturation, default value 0).

$$SI = \log(IAP/KT)$$

where,

IAP = the ion activity product for the given material and

KT = the reaction constant at the given temperature

If $SI > 0$, then the solution is super-saturated with respect to the mineral phase;

If $SI < 0$, then the solution is below saturation of the specified mineral phase.

If $SI = 0$, then the solution is in equilibrium with the specified mineral phase.

For Gas phases the SI corresponds to the log of the partial pressure.

Mols: Mols describes the maximum amount of mineral phase that can be added or dissolved to reach the specified saturation. The default value is 10 mols. The constraint on available amount of mols is important if you want to translate measured concentrations of minerals in an aquifer in a PHREEQC transport model. You may also set the amount of available mols to 0, making sure that the respective mineral can only be precipitated but not dissolved from the aquifer.

Alternative phase: The alternative phase is added or removed until the first mineral is in equilibrium with specified saturation index. For example, gypsum may be added or removed to the solution until an equilibrium with the primary mineral calcite is reached. An amount must be specified for the alternative mineral phase (default amount of the alternative phase is 10 moles as specified by PHREEQC). The solutions equilibration with calcite will be terminated if the amount specified for the alternative mineral phase is consumed prior to equilibration with calcite. It should be noted that if the alternative phase is specified, the amount of the mineral phase is ignored. Otherwise the system would be overdetermined.

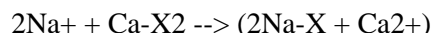
Below the grid, select the solution with which the mineral phase will be equilibrated. If you do not choose a solution, the specified assemblage is equilibrated with the first solution defined in the current simulation.

The equilibration of a solution with minerals will result in a new composition of the solution. The **Save Solution as** allows you to save this composition as a new solution. These new 'elements' can then be used as input data for subsequent PHREEQC simulations. Each of these new elements must be identified by a unique number that is not already in use. If you specify a number that is currently in use, the new element will overwrite the old element.

7.3.3 Exchange Assemblage

Ion exchange involves the replacement of one chemical in the solution with another on a solid surface. Intrusion of fresh water into marine sediments or seawater into coastal fresh water aquifers are probably the best known examples for this mechanism. Fresh water typically is dominated by Ca^{2+} and HCO_3^- ions due to the presence of dissolved calcite, whereas in seawater, Na^+ and Cl^- are the dominant ions. Sediment in contact with sea water will often have large concentrations of Na^+ .

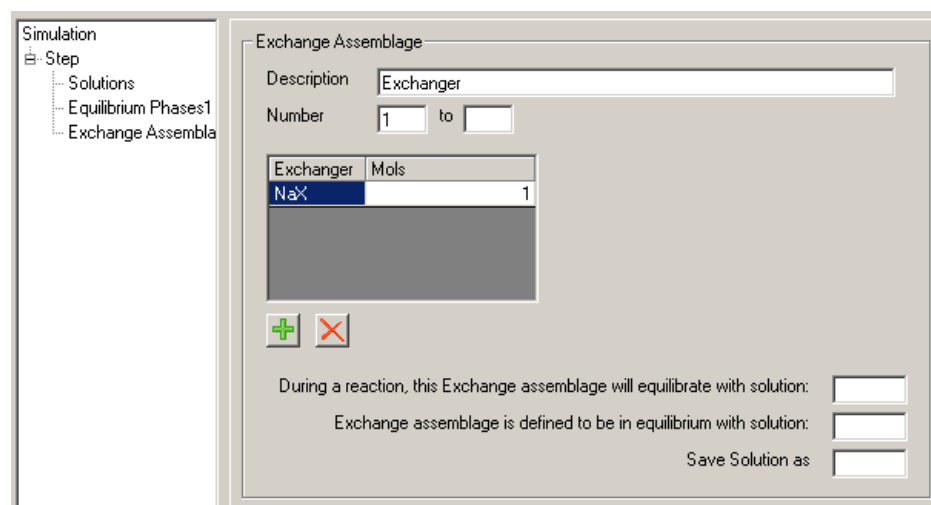
When seawater intrudes in a coastal fresh water aquifer an exchange of cations takes place:



where X indicates the soil exchanger.

Sodium is taken up by the exchanger and Ca^{2+} is then released. The composition of the solution and the exchanger is modified by this reaction.

A sample of the **Exchange Assemblage** options dialogue is seen below:





PHREEQC lets you define the initial composition of an **Exchange Assemblage** in two ways:


- [1] Explicitly by selecting the composition of the Exchange Assemblage. This approach is chosen if the mineralogy and the exchange sites for these minerals are well known.
- [2] Implicitly by specifying that the Exchange Assemblage is in equilibrium with a solution of a fixed composition. This approach is chosen if the type of water in the aquifer is known. E.g. to model an aquifer in contact with seawater make sure that your model contains a seawater sample and specify this sample within the “Exchange assemblage is defined to be in equilibrium with solution” option.

Option 1

AquaChem provides a list of available exchanger ions to choose from and allows you to easily add several ions to the Exchange Assemblage.

To add an ion(s) to the Exchange Assemblage:

- Click  to add a new line to the grid.
- Under the **Exchanger** field, double-click in the first empty cell and a combo box with a  button will appear listing the available items.
- Select the ion you need, and press **<Enter>** (on your keyboard). The ion will appear in the grid.
- Enter the Amount (in **Mols**) for each ion in the **Exchanger** column. In order to estimate the amount of exchangeable cations, you may use the **Aquifer Properties Calculator** described above.

To remove an ion from the Exchange Assemblage, click-on the ion you would like to remove, and then press the  button.

Option 2

You may specify that the Exchange assemblage is defined to be in equilibrium with a solution. For example, if you consider that the exchange takes place on a marine clay, define seawater as a solution and select this solution in the **Implicit** option as the water the exchange material is in equilibrium with. Simply enter a solution number in the appropriate field.

Once you have defined the Exchange Assemblage, you must select the solution with which you wish to equilibrate. If you do not select a solution from the list, you can save the Exchange Assemblage to equilibrate in a later simulation step (e.g. during a transport problem).

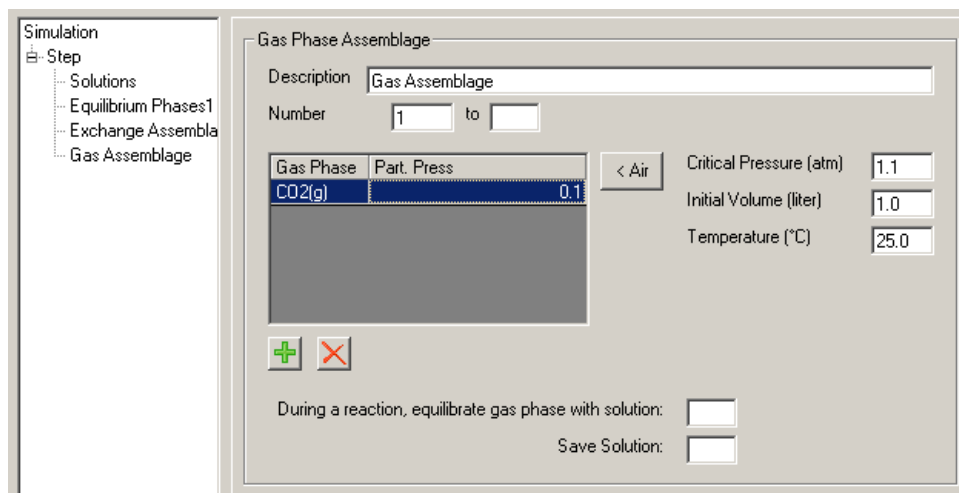
Each Exchange Assemblage is identified by a unique **Number** and a **Description** so they can be easily recognized in the Input / Output files, and reused in later simulations.

The **Save Solution as** field allows you to save the composition of a simulation by creating a new exchange assemblage. These new 'elements' can then be used as input data for subsequent PHREEQC simulations. Each of these new elements must be identified by a unique number that is not already in use. If you specify a number that is currently in use, the new element will overwrite the old element.

7.3.4 Gas Phase Assemblage

The Gas Phase is used to equilibrate a solution with a finite volume of gas. This simulation is not applicable if fixed partial pressures of the gas components are desired (for this case use the Mineral Phase option instead and fix a gas with a partial pressure

that has an infinite volume of gas phase). An example of the **Gas Phase Assemblage** dialogue is shown below:



The **Gas Phase** option will equilibrate the solution with only those gases which are defined in the simulation. However, gases may be specified with a partial pressure equal to zero at the start of the simulation. In this case, no moles of that component will be present initially, but some gases may evolve during a reaction simulation such as oxidation of organic material. As a result, the partial pressure of all gases included in the gas phase stays constant and the excess gas is removed from the solution as the gas bubbles.

Pressure calculations are computed using the ideal gas law,

$$PV = nRT$$

where,

P = defined pressure

V = volume



n = number of moles of the gas


R = universal gas constant

T = temperature

AquaChem provides a list of available gases to choose from and allows you to easily add multiple gases to the **Gas Phase Assemblage**.

To add a gas phase assemblage to the simulation:

- Click  to add a new line to the grid.
- Under the **Gas Phase** field, double-click in the first empty cell and a combo box with a  button will appear listing the available items.
- Select the gas you need, and press **<Enter>** (on your keyboard). The gas will appear in the grid.
- Enter a partial pressure value for each gas under the **Part. Press** column.

If you wish to add air to the solution, click on  button and the major components of air will be automatically added to the list, with their partial pressures at 1 bar.

To remove a gas from the **Gas Phase Assemblage** table, select the gas to remove and then press the  button.

The **Critical Pressure**, **Initial Volume** and **Temperature** information is required for each Gas Phase assemblage in order to calculate the weights of each gas component.

The default values are:

Critical Pressure: **1.1** atm. This is equal to the pressure, at which the dissolved gas will start to create bubbles

Initial Volume: **1.0** liters: amount of gas initially in the system

Temperature: **25°C**

Finally, you can select the solution with which you would like to equilibrate the gas phase assemblage. If you do not select a solution from the list, you can save the gas phase to equilibrate in a later simulation step. Each Gas Phase Assemblage is identified by a unique number and can be reused in later simulations.

The **Save Solution as** allows you to save the composition of a simulation by creating a new gas phase assemblage. These new ‘elements’ can then be used as input data for subsequent PHREEQC simulations. Each of these new elements must be identified by a

unique number that is not already in use. If you specify a number that is currently in use, the new element will overwrite the old element.

7.3.5 Adding Solutions

In PHREEQC, a "solution" is a set of analyses for a single water sample. AquaChem automatically creates a PHREEQC input file from the sample details pertaining to your AquaChem database, or from a pure water sample.

To access the solutions options, expand the items in the **Navigator tree** by pressing the **+** beside **Step**. Then select **Solutions** and the solutions frame should appear, similar to the one below.

The screenshot shows the AquaChem software interface. At the top, there is a table titled "Active Solutions in database". Below this table is a green plus icon and a button labeled "Add pure Water". Underneath, there is a section titled "Solutions" which contains a table with two columns: "Number" and "Description". At the bottom of the interface, there is a "Default Unit:" label followed by a dropdown menu set to "ppm".

ID	SAMPLEID	WATERTYPE	SAMPLE_DATE	STATIONID	SYI
4	MW-1-95	Ca-Na-SO4-Cl-HCO3	7/30/1995	MW-1	
5	MW-1-96	Ca-Na-SO4-Cl-HCO3	7/28/1996	MW-1	
6	MW-1-97	Ca-Na-SO4-Cl-HCO3	6/15/1997	MW-1	
7	MW-1-98	Na-SO4-Cl-HCO3	8/1/1998	MW-1	
8	MW-3-92	Ca-SO4-HCO3	8/8/1992	MW-3	
9	MW-3-93	Ca-HCO3-SO4	6/8/1993	MW-3	
10	MW-3-94	Ca-SO4-HCO3	6/15/1994	MW-3	

Number	Description
8	MW-3, 8/8/1992
2	pure Water


Default Unit: ppm


You must specify at least one solution for every PHREEQC simulation file. AquaChem allows you to select from two types of solutions:

- Samples from your AquaChem database; OR
- Pure Water

These options are described in detail in the following sections.

Using Samples from your AquaChem Database


To add a sample from your database as a solution, select the sample from the list at the top, and press the  button. The sample will then be added to the **Solutions** frame in the lower half of the dialogue; in addition, the new solution will show up as a new line in the **Navigation tree**, under **Solutions**.

To add multiple samples from your database, use the <Shift> and <Ctrl> keys with the mouse to select the samples you need, then click the  button. This is useful if you want to speciate a large number of samples simultaneously. Each complete sample should have pH, alkalinity, and temperature values specified.

The AquaChem database index **Number** for the selected sample will be used as the Solution number. The **Description** field is used to identify the solutions. The solution description will appear in the PHREEQC input file.

After you have added a solution, you can verify that it has been added to the PHREEQC input file. To do so, click the **Input File** tab at the bottom of the screen and you should see the added solution and any corresponding concentrations.

Adding Pure Water

To add pure water as a solution, press the  button below the list of AquaChem active solutions. For certain simulations it might be useful to have pure water as a base solution. For example, you can simulate a groundwater by saturating pure water with calcite and adding a specified CO₂ pressure.

Solution Properties

Once you have added a solution, you may want to modify the Solution properties.

To do so, you have two options:

- In the **Solutions** frame, double click on the appropriate solution from the main dialogue, OR
- Select the appropriate **Solution #**, from the **Navigator tree** on the left side of your display.

The **Solutions** properties frame should appear as shown below.

The screenshot shows the 'Solution' dialog box with the following details:

- Description:** MW-3, 8/8/1992
- Number:** 8 to []
- General Tab:**
 - pH:**
 - ☒ Constant during reaction: 7.4
 - ☐ Determined by equilibration with phase: [] [0]
 - ☐ Charge (not recommended)
 - Redox:**
 - ☒ Constant during reaction: []
 - ☐ Determined by equilibration with phase: [] []
 - ☐ Charge (not recommended)
 - ☐ Redox Couple: []
- Temperature (°C):** 14
- Density (g/cm3):** []

The upper part of this dialogue lists the solution **Description** and **Number**. As discussed above, the **Number** is AquaChem database index number for the selected sample which will be used as the Solution number. The **Description** field is used to identify the solutions.

The remaining options in this frame are separated into two tabs:

- **General**, and
- **Concentrations**

The options associated with these tabs are described in detail in the following section.

General

Under the **General** tab, you can specify options for **pH** and **Redox** during the simulation.

You can choose a **Constant during reaction** value for these parameters, or allow the values to be determined as a result of equilibration with a mineral phase (**Determined by equilibration with phase**).

In addition, at the bottom of the frame, you can enter a default **Temperature** and **Density** values for the solution. The sample density is only needed when concentration values are entered in ppm or ppb.

Concentrations

This tab lists the concentrations of the dissolved elements in solution.


Element	M. Spec.	Conc.	Unit	Phase	Sat.
H	H+				
H(0)	H2				
H(1)	H+				
E	e-				
O	H2O				
O(0)	O2				
O(-2)	H2O				
Ca	Ca+2	256	mg/l		
Mg	Mg+2	19	mg/l		
Na	Na+	50	mg/l		
K	K+	2	mg/l		
Fe	Fe+2	0.3	mg/l		
Fe(+2)	Fe+2				
Fe(+3)	Fe+3				
Mn	Mn+2	0.29	mg/l		

Default Unit: mg/l

The first column of the grid contains the **Element** name, while the second column describes the PHREEQC Masterspecies (**M. Spec.**) name.

The parameter values are read from the database; if necessary you may add or remove concentrations (**Conc.**), or edit the existing values. You may also define a concentration as controlled by a mineral solubility, or for gases, by a defined pressure. For example, a calcium concentration is overridden in the simulation by a calculated value that will result in a calcite equilibrium state (Saturation index = 0).


Once you have entered element values for the pure water solution, the **Default Unit** may not be changed. However, you can change the unit individually for each entered

value by clicking  button in the **Unit** column. Please note that it is not possible to freely mix units (i.e. you may not enter some values as mg/L and others as mmol/l). You may only mix units within a "unit family" such as g/L, mg/L ug/L.

Example

The following example will demonstrate how to use a sample from the Demo database, as a solution in a PHREEQC simulation:

To add the **MW-1-92** sample from the **DEMO.AQC** database to a PHREEQC input file, proceed as follows:

- **Tools** from the main menu, then select **Modeling** and then **PHREEQC (Basic)** from the main menu.
- Click **+** beside **Step** (in the **Navigator tree**) and you will see **Solutions**
- Select **Solutions**
- Select the **MW-1-92** sample (ID = 1) at the top of the list of **Active solutions in database** frame
- Click  button and the sample will now appear in the **Solutions** frame in the bottom section of the dialogue.

To view the solution properties, double click on this **Solutions**, or expand the **Navigator tree**, and select the **Solution** from here. This will load a **Solution** properties dialogue.

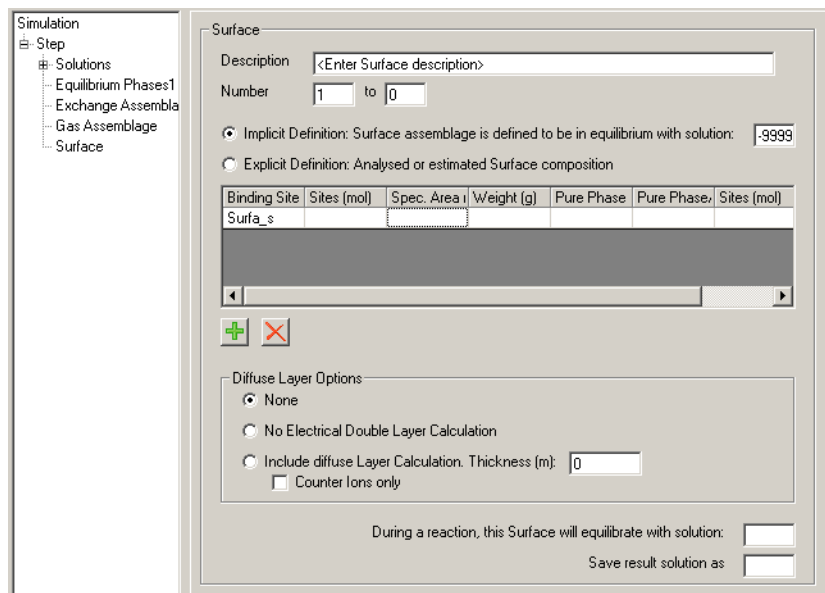
Click on the **Concentrations** tab to confirm that the concentrations are identical to those entered for that sample in the AquaChem database.

To verify that this data is saved to the PHREEQC input file, you can view the input file by clicking the **Input File** tab at the bottom of the dialogue.

7.3.6 Surface Assemblage

While ions are exchanged during an exchange process, surface processes involve only the absorption of material on mineral surfaces. This process typically can be observed on hydroxides and is driven by electrostatic processes.

An example of the **Surface** options dialogue is seen below:





PHREEQC lets you define the initial composition of a **Surface Assemblage** in two ways:

1. Explicitly by specifying the parameter values for Surface Assemblage; or
2. Implicitly by specifying that the Surface Assemblage is in equilibrium with a solution of a fixed composition.

Option 1 (Explicit)

To define a Surface Assemblage:

- Click  to add a new line to the grid.
- Under the **Binding Site** field, double-click in the first empty cell and a combo box with a  button will appear listing the available items.
- Select a Binding site, and press **<Enter>** (on your keyboard).
- Overwrite the default values for **Specific Area** (600 m²) and **Weight** (89 g) if required. Weight is defined as mass of solid for calculation of surface area in grams. Surface area is mass times specific area per gram.
- Specify the total number of **Sites** (in **mols**) for each binding and make any appropriate changes to the default values for **Specific Area** and **Weight**.

- Choose the appropriate diffuse layer model for the surface reaction (please see the PHREEQC manual for information on selecting the appropriate **Diffuse Layer Options** for your surface reactions).

To remove a binding site from the Surface Assemblage, select the **Binding Site** you would like to remove and press the  button.

Option 2 (Implicit)

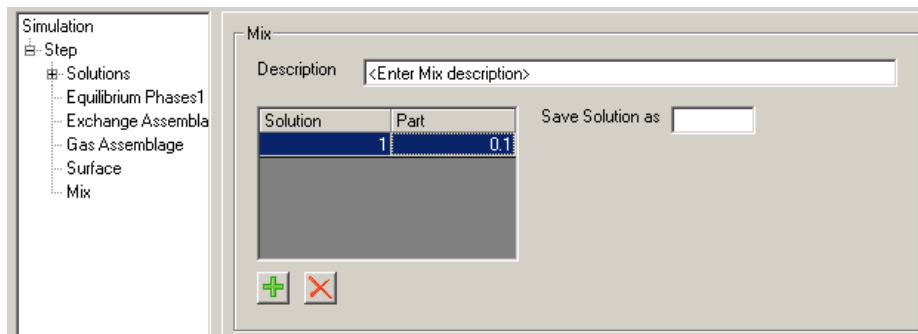
You may specify that the Surface Assemblage is defined to be in equilibrium with a solution. Simply enter a solution number in the appropriate field.

Once you have defined the Surface Assemblage, you must select the solution with which you wish to equilibrate. Each Surface Assemblage is identified by a unique **Number** and a **Description** so they can be easily recognized and reused in later simulations.

The **Save Solution as** allows you to save the composition of a simulation by creating a new surface assemblage. These new ‘elements’ can then be used as input data for subsequent PHREEQC simulations. Each of these new elements must be identified by a unique number that is not already in use. If you specify a number that is currently in use, the new element will overwrite the old element.


7.3.7 Mix Solutions

The Mix Solutions allows you to simulate the mixing of multiple solutions together with different proportions of each solution added to the mixture. An example of the **Mix** options dialogue is seen below:



To simulate mixing multiple samples together, you must first specify at least two solutions for the PHREEQC simulation.

To add a solution to the mixture:

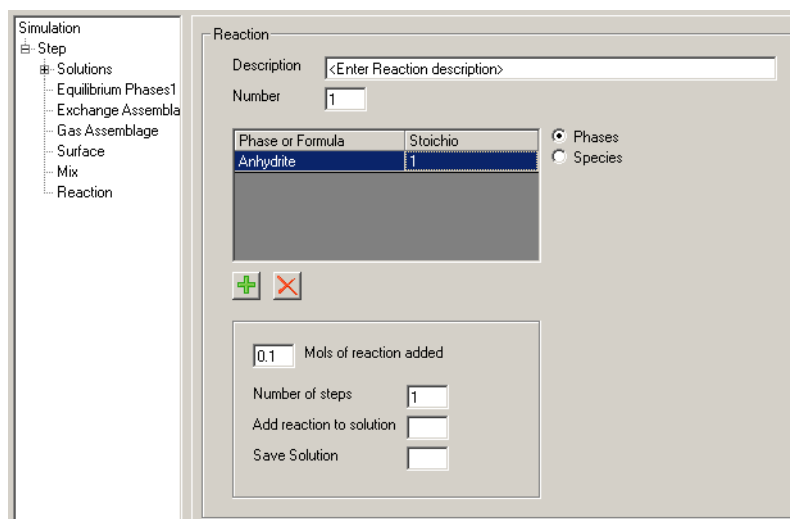
- Click  to add a new line to the grid under the **Solution** header.
- Type in a Solution number in the same cell.
- Click-on the first line in the grid under the **Part** header and type the proportion of the solution you would like to add to the mixture (e.g. 3 parts of Solution 1 and 4 parts of Solution 2)

The solutions will be automatically added to the input file. Note that if you want to simulate several mixing proportions between two samples, each mix must be conducted in a separate simulation step as otherwise only the first mix definition will be executed.

Each mixed solution has a unique Mix number and **Description**. To re-use the simulated mixed sample in a later step, simply enter a solution number beside the **Save Solution as** field.

7.3.8 Reactions

The Reactions simulation options are used to dissolve or precipitate a specified amount of a mineral or formula. To access these options, you need to add a **Reaction** as a **Step** and then access the **Reactions** options. Below is an example of the **Reaction** dialogue:



AquaChem provides a comprehensive list of available reactant species and phases to choose from. To select from the list of available reactants:

- Specify to show **Phases** or **Species**
- Click to add a new Phase or Species. A new line will be added to the grid.
- Under the **Phase or Formula** field, double-click in the first empty cell and a combo box with a button will appear listing the available reactants.
- Select the reactant you need, and press **<Enter>** (on your keyboard). The reactant will appear in the grid.
- Enter the stoichiometric factor in the **Stoichio** field beside the reactant. The stoichiometric factor defines whether the reactants are added ($SF > 0$) or removed ($SF < 0$) from the solution. The stoichiometric factor is multiplied by the Moles of reaction added in order to calculate the total amount of the respective reactant added (default value is **1**).
- Type the number of **Moles of reactant added**, the **Number of steps** for the reaction, and **Add reaction to solution** number. The reaction path is revealed when you simulate the reaction in multiple steps. By default the reaction is added to the first solution in the simulation. Note that you are not required to add the reaction immediately to a solution if you only wish to use the reaction in later simulation step(s).
- To add additional reactions, you must add additional Reaction steps. To do this, select **Reactions** from the **Navigator tree**. Then, under the **Reactions** frame, click to add a new Reaction. A new line will be added to the navigator tree. Select this new item, and the options will be loaded for this new reaction.

The **Save Solution as** allows you to save the composition of a simulation by creating a new reaction. These new ‘elements’ can then be used as input data for subsequent PHREEQC simulations. Each of these new elements must be identified by a unique number that is not already in use. If you specify a number that is currently in use, the new element will overwrite the old element.

The reaction will be added automatically to the input file. To verify this, click the **Input File** tab at the bottom of the dialogue.

NOTE: The add reaction option does not calculate an equilibrium status. It will only dissolve the specified amount of chemicals in the given solution. Therefore you may enter any valid chemical formula into the phase or formula grid

The following example demonstrates how to add a reaction.


Example

Follow the steps listed below to specify the following reaction:



Dissolve 0.1 moles **Calcite** and 2.0 moles **Halite** (NaCl) in pure water.

- Select **Tools** from the main menu, then **Modeling** and then **PHREEQC (Basic)** from the main menu.

The **PHREEQC Input** dialogue will appear, and the **General** tab will be active.



- Click **+** beside **Step** (in the **Navigator tree**) and you will see **Solutions**
- Select **Solutions**
- Add **pure water** as a solution by clicking  below the active solutions frame
- Click **Step** from the **Navigator tree** and right click on it
- Select **Add Forward Model** and then select **Add Reaction** to add the First Reaction step.
- **Reactions** a new item **Reaction1** will now appear in the **Navigator tree**.

Under the **Reaction** options (in the **Navigator tree**):

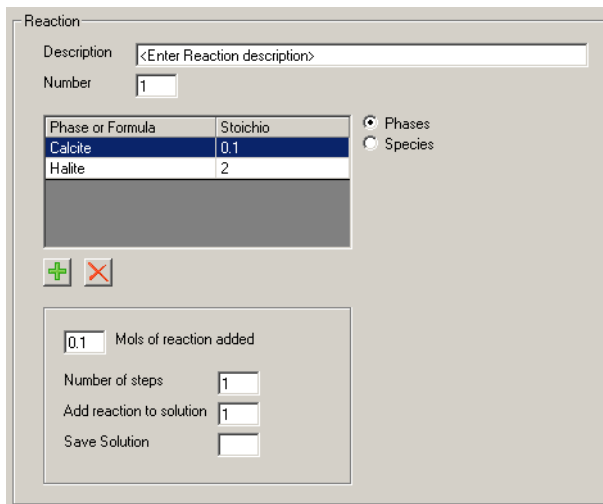
- Select **Reaction1** in the **Navigator tree**
- **Phases** radio button, under the **Reaction** options frame
- Click  to add a new Phase or Species. A new line will be added to the grid.
- Under the **Phase or Formula** field, double-click in the first empty cell and a combo box with a  button will appear listing the available reactants.
- Select **Calcite**, overwrite the default value of **1.0** with **0.1**. Press **<Enter>** on your keyboard.

This reaction will be automatically added to the input file. To verify this, click the **Input File** tab at the bottom of the dialogue.

To add a reaction for **Halite**:

- Click  to add a new Phase or Species. A new line will be added to the grid.
- Under the **Phase or Formula** field, double-click in the first empty cell and a combo box with a  button will appear listing the available reactants.
- Select **Halite**, and a default value of **1.0** will show. Press **<Enter>** on your keyboard.
- type: **1** in the **Mols of reaction added** field (located below the grid)
- Select the field labeled **Add reaction to solution**.
- Type **1** in this field.

Once you have added this reaction, your dialogue should appear similar to the one shown below:





Reaction

Description:

Number:

Phase or Formula	Stoichio
Calcite	0.1
Halite	2

☒ Phases
☐ Species

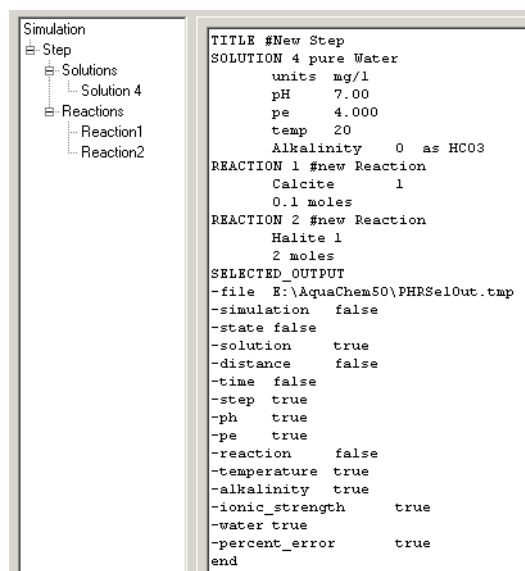
Mols of reaction added

Number of steps:

Add reaction to solution: ☒

Save Solution: ☐

To verify the input file, click on the **Input File** tab at the bottom of this dialogue. The contents of the input file should be similar to that shown in the figure below:



You may now run PHREEQC with these reaction and solution options.

7.4 Running PHREEQC Simulation

Once you have specified the necessary Solutions and Reaction Steps, you are ready to run the simulation. Before doing so, it is recommended that you verify the contents of the input file. Once this is finished, click the **[Run]** button to run the simulation.

The PHREEQC simulation will run in the background as a DOS application inside a DOS window. Once the simulation is finished, you can verify the results by clicking the **Output File** tab (at the bottom of the main PHREEQC dialogue) and scan this for any error messages. The output from the PHREEQC run can be viewed by clicking on the **Output File** tab, or on the **Spreadsheet** tab.

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Appendices

Appendix A: Troubleshooting and FAQ

The following list provides answers to some of the most common problems encountered by users of AquaChem. Please review this list of frequently asked questions before contacting SWS Technical Support for assistance with your project.

Q 1: *My sample symbols do not show up on the plot.*

- A1. Check that you have assigned symbols to that sample(s).
- A2. Ensure the symbol group is active for the plots. This can be set in the **Plots > Define Symbol or Line** dialogue.
- A3. Ensure that you have entered data for each parameter required by that plot.

Q 2: *My symbols show up on the plot colored in red, even though I have assigned another color to this symbol.*

- A. The red colored symbols correspond to the samples which are selected in the active list. This indicates you are using the **Identify Samples** option in the **Define Symbol or Line** dialogue. Disable this option and the symbols will no longer be identified.

Q 3: *After I import data, the data does not appear correctly or does not appear at all.*

- A1. Check that you have a value for each sample for each parameter in your source file.
- A2. Ensure that you have selected the correct **Delimiter** during the import routine.
- A3. Check your data source file to ensure that there are no extra spaces between columns or rows of data.

Q 4: *During data import, I get an error “Column must be assigned to CAS Registry #.”*

- A1. Ensure that you have specified the correct import format. For this warning, you should typically try the **Samples as Rows** in the **Format** field.
- A2. If your data source file does not contain CAS Registry #'s, ensure that this option is disabled during the data import.

Q 5: *During data import, I get error “Column must be assigned to SampleID or StationID.”*

- A1. Ensure that you have specified the correct import format. For this warning, you should typically try the **Samples as Rows** in the **Format** field.
- A2. Ensure that you have mapped at least one of your text fields as the StationID and SampleID.

Q 6: *Some of my samples or stations no longer appear in the active list.*

A. Since samples and stations are actively linked in the database, when you temporarily omit a station, the corresponding samples will be removed from the active list. Likewise, when you remove all samples assigned to a given station, this station will be temporarily removed from the active list. To restore all samples or stations press the **Show all** button in the active samples list dialogue, or select **Filter** from the main menu and then **Show all** option.

Q 7: *In the print options window, part of the plot is cut off by the print template.*

A. Be sure to select the print template first, then load the plot on top of the template. This will ensure the page margins are set correctly, and will not cut off sections of the plot.

Q 8: *After I enter a value for a measured parameter, the 0 after the decimal place is cut off (for example, 7.20 is entered, however AquaChem displays this as 7.2).*

A. This can be corrected by modifying the parameter format. Go to File > Database > Parameters. Select the appropriate measured parameter, and modify the format. Disable the Auto option, and increase the number of decimal places to the desired value.

Q 7: *Is it possible to create two unique Piper plots showing two different sample groups?*

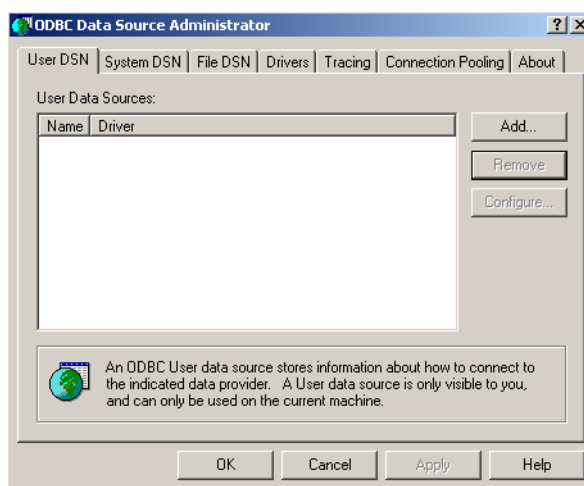
A. AquaChem will not allow for multiple Piper plot windows showing two distinct sample groups, at the same time. Since plots are created from information based on the active samples list, any changes that are made in the active list will immediately impact any open plot windows. The plots must be created individually.

Appendix B: Configuring an ODBC Connection

To use the AquaChem - HGA linking feature, you must configure an ODBC connection for your HGA SQL Server database. This can be done using Windows native ODBC Data Source Administrator tool. The following describes the steps to configure an ODBC connection:

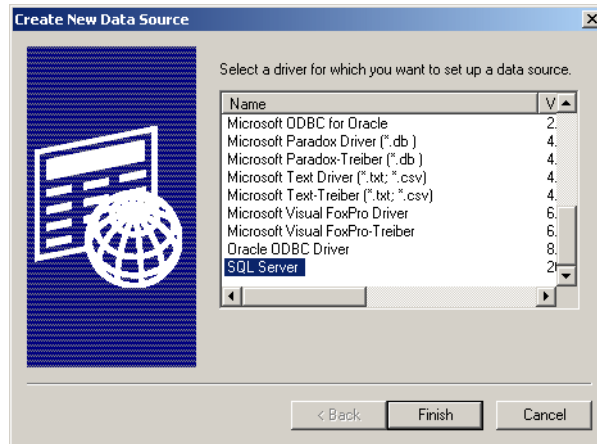
Please note that the following may vary slightly between different operating systems. The following steps are described for Windows XP.

1. Go to **Start / Settings / Control Panel / Administrative Tools**
2. Select **Data Sources (ODBC)**. The following windows dialog will display:

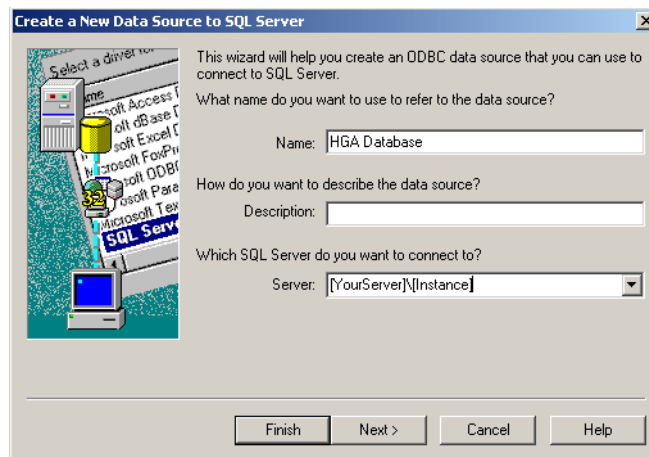


From the User DSN tab, click the **[Add]** button.

3. From the **Create New Data Source** dialog, select **SQL Server** and then click the **[Finish]** button



4. From the **Create a New Data Source to SQL Server** dialog, enter the following information, and then click **[Next]**.

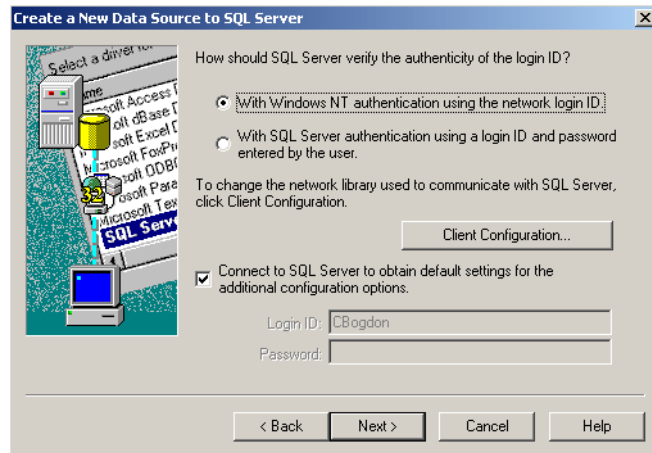


Name: Enter a name for the new data source

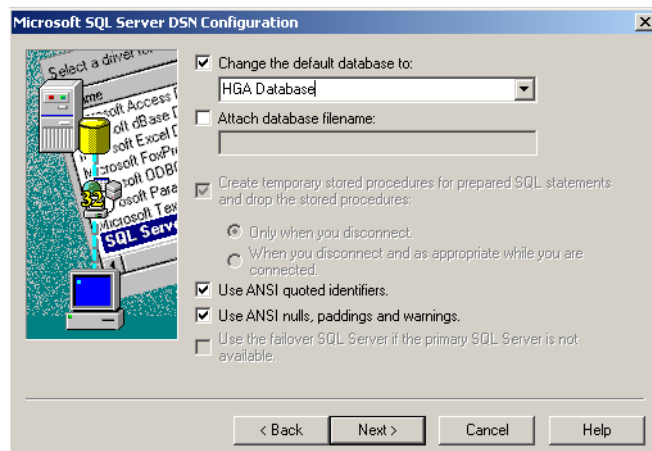
Description: Enter a description for the new data source (optional)

Server: Specify the server and instance that contains the HGA database. If the database resides on your local machine, then you would specify your computer name, instead of a server name.

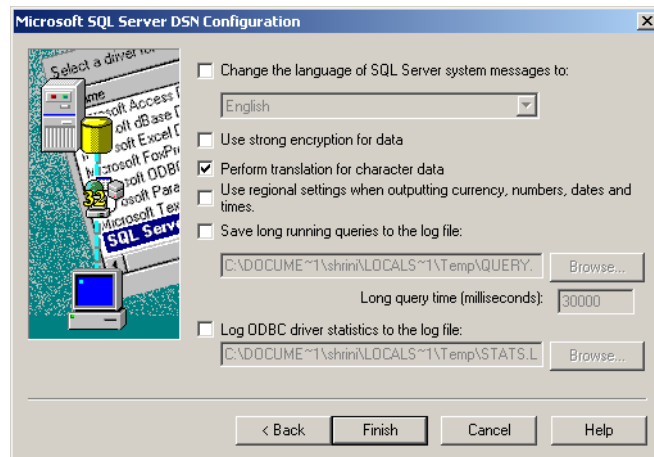
5. In the next dialog, select the appropriate SQL Server authentication option. In most cases, you can leave the default settings and then click the **[Next]** button.



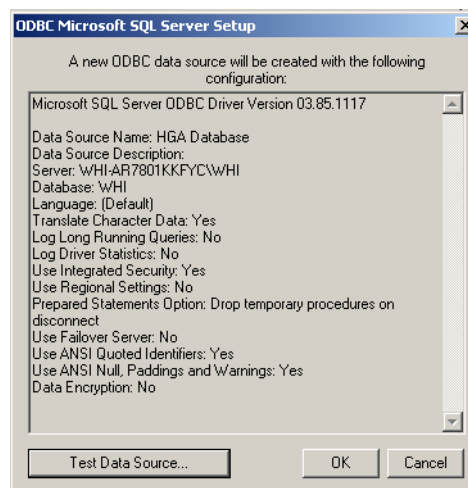
- Choose your HGA database from the **Change the default database to:** combobox and select **[Next]**.



7. In the next dialog, leave the default settings, and select the **[Finish]** button.



8. Finally, click the **[Test Data Source...]** button to test your ODBC connection.

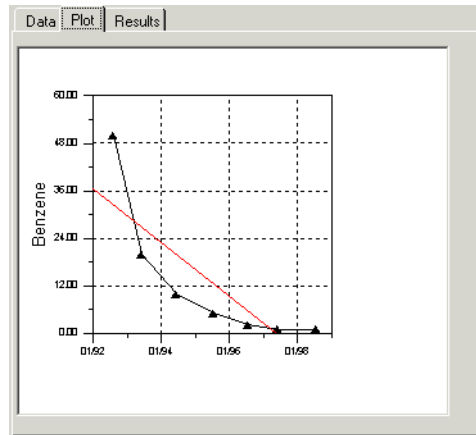


If successful, click **[Ok]** to finish. If the test failed, review the ODBC settings to ensure that they are correctly defined.

Appendix C: Trend Analysis Tests

Linear Regression

The linear regression simply calculates a regression line on the time/value plot. A positive slope of the regression line indicates a trend towards increasing values, a negative slope a trend to decreasing values. However the result of this test should only be used qualitatively and should be confirmed by more sophisticated tests such as Sen's and Mann-Kendall. The calculated line is drawn on the time series plot.



Sen's Test

Sen (1968) developed a simple nonparametric estimator of trend which is particularly useful for groundwater monitoring applications. The method is robust to outliers, missing data and non-detects. To compute the Sen's trend estimator S , the individual slopes for any succeeding points are calculated. The trend estimator corresponds to the median of all calculated slopes. To calculate the confidence limits, an estimate of the variance of S is required which may be calculated using the following formula:

$$\text{var}(S) = \frac{1}{18} \left[n(n-1)(2n+5) - \sum_{p=1}^q t_p(t_p-1)(2t_p+5) \right]$$

where q is the number of values for which there are ties and t_p is the number of tied measurements for a particular value. The lower and upper confidence limit M_1 and M_2 can be calculated as follows:

$$M_1 = \frac{N' - Z_{1-\alpha} \sqrt{\text{var}(S)}}{2}$$

$$M_2 = \frac{N' + Z_{1-\alpha} \sqrt{\text{var}(S)}}{2}$$

The calculated ranks M_1 and M_2 are projected on the ranked slope data. If the respective interval contains the zero value, the Null Hypothesis of non trend is accepted. The Output of Aquachem includes the Median slope, the variance S , M_1 , M_2 as well as their respective slopes. Both Hypothesis, “Increasing trend” and “Decreasing trend” are tested according to the specified probability and the result reported in the result table shown below.

The screenshot shows a software window titled "Trend analysis" with the following settings and results:

- Method:** Sen's test
- Parameter:** Ca mg/l
- Confidence:** 95% one sided

The "Results" tab is active, displaying a table with the following data:

Variable	Value	Comment
n	30	Number of datapoints
N'	435	Number of slopes
S	-0.50	Median slope (Sen's slope estimate)
var(S)	3141.667	variance of S
M1	171.3984	Rank M1
Q(171.3984)	-3.3333	Lower confidence level of slopes
M2	263.6016	Rank M2
Q(263.6016)	1.00	Upper confidence level of slopes
Result	Hypothesis of increasing trend rejected Hypothesis of decreasing trend rejected	

Buttons at the bottom include "Copy", "Calculate", and "Cancel".

For more details on the Sen's test please see the reference below:

Gibbons, R.D., 1995: *Statistical Methods for Groundwater Monitoring*. p.175.

Mann-Kendall Test

The **Mann-Kendall** test is a trend estimator that can be used to prove if contaminant concentrations are significantly diminishing or rising over time. For this test, there are no distributional assumptions, and missing data (non-detects) or irregularly spaced measurement periods are allowed. Non-detects are assigned a value smaller than the smallest measured value.

The version of the **Mann-Kendall** Test used in AquaChem can be applied for virtually any groundwater parameter. The **Mann Kendall** test provides two values; S value and Z value. The test procedure is as follows:

First, order the data by sampling date x_1, x_2, \dots, x_n where x_i is the measured value on occasion i .

Second, record the signs of each of the N' possible differences $x_{i'} - x_i$ where $i' > i$. For example, let

$$\begin{aligned}\text{sgn}(x_{i'} - x_i) &= 1 \text{ if } x_{i'} - x_i > 0 \\ &= 0 \text{ if } x_{i'} - x_i = 0 \\ &= -1 \text{ if } x_{i'} - x_i < 0\end{aligned}$$

The Mann-Kendall statistic (S) is then computed as:

$$S = \sum_{i=1}^{n-1} \sum_{i'=i+1}^n \text{sgn}(x_{i'} - x_i)$$

which is the number of positive differences minus the number of negative differences.

If $S = 0$, then there is no increasing or decreasing trend in the data

If $S < 0$, then there is a decreasing trend, indicating concentration is decreasing over the time interval.

If $S > 0$, then there is an increasing trend, indicating concentration is increasing over the time interval.

A two-sided test (for either increasing or decreasing trend) can also be obtained, using probability values. For $n > 10$, then the normal approximation (Z) is calculated as follows:

If $S > 0$, then

$$Z = \frac{S - 1}{[\text{var}(S)]^{0.5}}$$

If $S = 0$, then $Z = 0$

If $S < 0$, then

$$Z = \frac{S + 1}{[\text{var}(S)]^{0.5}}$$

Where $\text{var}(S)$ is calculated according to the formula used for the Sen's slope estimator. The quantity Z can be compared to standard normal cumulative distribution probabilities to test the null hypothesis of no trend.

In the **Statistics Report** window, the S value is displayed as $S(M-K)$, and the Z value is displayed as $Z(M-K)$.

For more details on the Mann-Kendall test please see the reference below:

Gibbons, R.D., 1995: *Statistical Methods for Groundwater Monitoring*. p.178.

After you have selected a test to use in trend calculation, select the parameter for which you wish to calculate the trend.

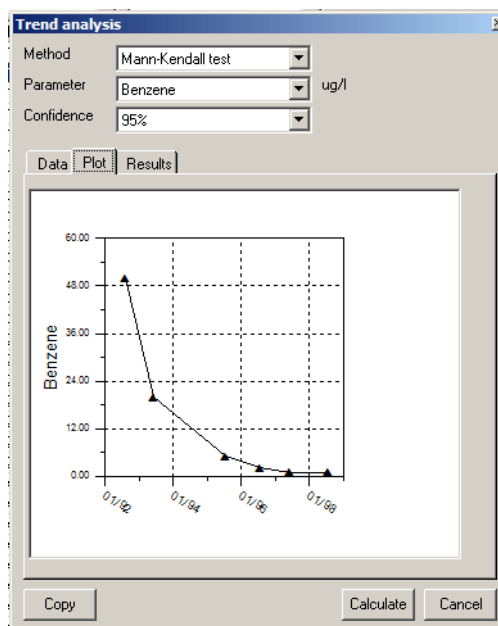
All stations will be displayed in the **Data** tab, sorted by date in ascending order. At this point you can de-select any samples you do not wish to use in the analysis. Simply uncheck the box beside the sample to remove it from calculations.

The screenshot shows the 'Trend analysis' dialog box with the 'Data' tab selected. The 'Method' is set to 'Mann-Kendall test', the 'Parameter' is 'Benzene' (with units 'ug/l'), and the 'Confidence' is '95%'. The 'Raw Data' table is displayed below, showing columns for 'Use', 'SampleID', 'Benzene', 'Date', and 'Delta'. The table contains 18 rows of data, with the first row highlighted in blue. The 'Use' column has checkboxes for each row, with the first, second, and third rows checked.

Use	SampleID	Benzene	Date	Delta
<input checked="" type="checkbox"/>	OW-4-92	0.5	7/15/1992	
<input checked="" type="checkbox"/>	OW-2-92	50	8/1/1992	17
<input checked="" type="checkbox"/>	MW-3-92	0	8/8/1992	7
<input type="checkbox"/>	MW-1-92	50	8/15/1992	7
<input type="checkbox"/>	OW-4-93	0.5	5/25/1993	283
<input type="checkbox"/>	MW-1-93	20	6/1/1993	7
<input checked="" type="checkbox"/>	OW-2-93	20	6/5/1993	4
<input type="checkbox"/>	MW-3-93	0	6/8/1993	3
<input type="checkbox"/>	OW-4-94	0.5	5/23/1994	349
<input checked="" type="checkbox"/>	OW-2-94	10	6/12/1994	20
<input type="checkbox"/>	MW-1-94	10	6/15/1994	3
<input type="checkbox"/>	MW-3-94	0	6/15/1994	0
<input type="checkbox"/>	OW-4-95	0.5	6/12/1995	362
<input checked="" type="checkbox"/>	OW-2-95	5	7/21/1995	39
<input checked="" type="checkbox"/>	MW-3-95	0	7/25/1995	4

At the bottom of the dialog box, there are buttons for 'Copy', 'Calculate', and 'Cancel'.

Once you are satisfied with the data selection, move on to the **Plot** tab. This tab offers a graphical representation of the selected data.



The graphical representation of the data is very helpful because it allows you to see if there are any outliers that may potentially distort the analysis.

Once you are satisfied with the view click the **[Calculate]** button, and the trend will be calculated and displayed in the result tab.

The 'Trend analysis' dialog box is shown with the 'Results' tab selected. The table below contains the calculated statistics and the final result of the trend analysis.

Variable	Value	Comment
n	6	Number of datapoints
S	-14.00	Mann Kendall test statistic
Var(S)	27.3333	Variance
Z, z(1-a)	4865, 1.645	Normal approximation
Probability	-0.1376	Probability of no trend
Result	Hypothesis of increasing trend rejected Hypothesis of decreasing trend accepted	

Depending on the chosen test, different results will be displayed, however one common entry is the **Result** line at the bottom of the list of calculated parameters. The result can be:

- increasing trend accepted/rejected
- decreasing trend accepted/rejected

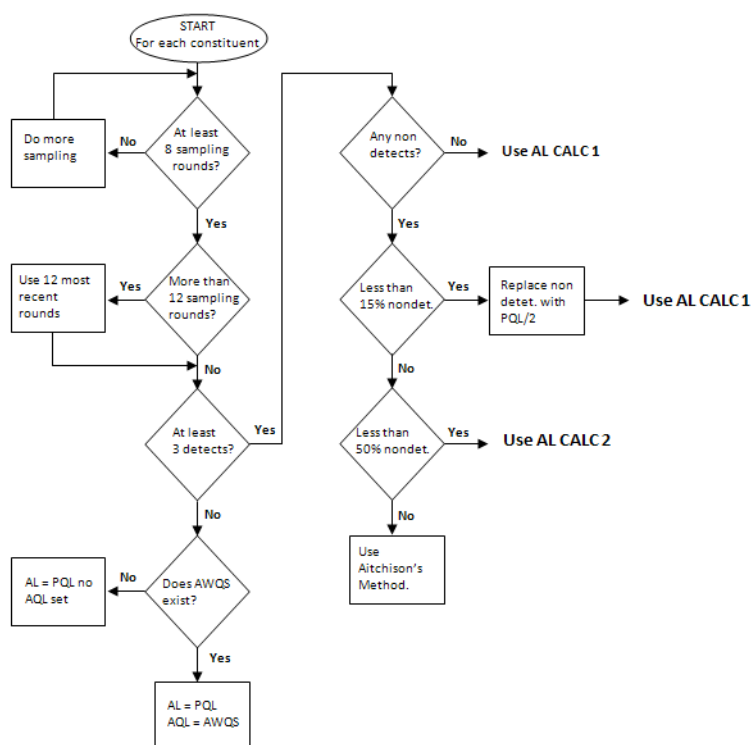
To see the calculations for another method and/or parameter simply change these options in their respective combo boxes and click **[Calculate]** again.

Appendix D: Calculating Alert Levels

Tolerance Interval Method

The tolerance interval method is one statistical method used in AquaChem to calculate upper and lower tolerance limits for groundwater constituents. Tolerance limits, or alert levels, are derived from historical compliance groundwater monitoring data. They may be useful for comparing with future monitoring data for detecting potential groundwater contamination. Please note that this method assumes that the historical data is of normal distribution.

Depending on the number of non-detects present in the historical dataset, one of three formulas are used to determine the appropriate alert levels: **Basic Tolerance Interval Method**, **Cohen's Method** or **Aitchison's Method**. The latter two methods are intended to handle datasets with a larger percentage of non-detects. The algorithm for determining which method to use is presented graphically below. All acronyms are described at the end of the appendix.



The formulas for each method is described on the following page.

AL Calc 1 : Basic Tolerance Interval Method

$$AL = \bar{x} + Ka$$

$$\bar{x} = \text{Mean} = \frac{\sum_{i=1}^n x_i}{n}$$

K = Tolerance factor

$$a = \text{std. dev.} = \sqrt{\frac{\sum_{i=1}^n (x_i - \bar{x})^2 n_i}{(n-1)}}$$

n = total number of quantifiable samples

AL Calc 2 : Cohen's Method

$$AL = \bar{x}_{corr} + Ka_{corr}$$

x = mean = as in AL Calc1 (see above)

K = tolerance factor from table 1

a = std. dev = as in AL Calc1 (see above)

n = total number of quantifiable samples

m = total number of samples

$m - n$ = total number of non detects

$h = (m-n)/m$ = corr. factor 1

$$g = \frac{a^2}{(\bar{x} - PQL)^2} = \text{corr. factor 2}$$

Obtain e from *Correction for Non-detects* table

$$\bar{x}_{corr} = \bar{x} - e(\bar{x} - PQL)$$

$$a_{corr} = \sqrt{a^2 + e(\bar{x} - PQL)^s}$$

Description of Acronyms

AL - Alert Level: an early warning indicator of a potential violation of a water quality standard at the applicable point of compliance.

AQL - Aquifer Quality Limit: a maximum concentration of a constituent allowed in an aquifer at the point of compliance. Synonymous with AWQS.

AWQS - Aquifer Water Quality Standards: numeric water quality standards

MCL - Maximum Contaminant Level: the maximum permissible level of a contaminant in water

MDL - Method Detection Limit: the minimum concentration of a constituent that can be measured and reported with a 99 percent confidence that the true value is greater than zero in a given matrix containing the analyte.

PQL - Practical Quantitation Limit: the lowest level that can be reliably achieved within specified limits of precision and accuracy during routine laboratory operating conditions.

Source: *Alert Level Guidance for Solid Waste Facilities - 1998*, Arizona Department of Environmental Quality.

